

This Page Is Inserted by IFW Operations  
and is not a part of the Official Record

## **BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

**IMAGES ARE BEST AVAILABLE COPY.**

**As rescanning documents *will not* correct images,  
please do not report the images to the  
Image Problem Mailbox.**

**THIS PAGE BLANK (4870)**

GenCore version 5.1.3  
Copyright (c) 1993 - 2002 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: November 19, 2002, 17:32:44 : Search time 86.2973 Seconds  
(without alignments)  
318.082 Million cell updates/sec

Title: PCT-US02-27145-2

Perfect score: 1048

Sequence: 1 MAEDADNMNELEEMGRADQ.....SNKTRIDEANQRATKMLGSG 206

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : A.Geneseq.101002.\*

```
1: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
2: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
3: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1982.DAT:*
4: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:*
5: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:*
6: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1985.DAT:*
7: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1986.DAT:*
8: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:*
9: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1988.DAT:*
10: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1989.DAT:*
11: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1990.DAT:*
12: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1991.DAT:*
13: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1992.DAT:*
14: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:*
15: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1994.DAT:*
16: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1995.DAT:*
17: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:*
18: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1997.DAT:*
19: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1998.DAT:*
20: /SID2/gcgdata/geneseq/geneseqp-emb1/AA1999.DAT:*
21: /SID2/gcgdata/geneseq/geneseqp-emb1/AA2000.DAT:*
22: /SID2/gcgdata/geneseq/geneseqp-emb1/AA2001.DAT:*
23: /SID2/gcgdata/geneseq/geneseqp-emb1/AA2002.DAT:*
```

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1048	100.0	206	18	AAW30103
2	1048	100.0	206	19	AAW79198
3	1048	100.0	206	19	AAW43426
4	1048	100.0	206	22	AAU00246
5	1048	100.0	206	22	AAU00253
6	1043	99.5	206	22	AAU02640
7	1042	99.4	206	22	AAU00259
8	1042	99.4	206	22	AAU00260
9	1042	99.4	206	22	AAU00261
10	1042	99.4	206	22	AAU02638

11	1041	99.3	206	22	AAU00262	Synaptosomal-assoc
12	1039	99.1	206	22	AAU00257	Synaptosomal-assoc
13	1039	99.0	206	22	AAU00266	Synaptosomal-assoc
14	1037	99.0	206	22	AAU02171	Synaptosomal-assoc
15	1036	98.9	206	22	AAU00256	Synaptosomal-assoc
16	1036	98.9	206	22	AAU02639	Synaptosomal-assoc
17	1033	98.6	206	22	AAU00258	Synaptosomal-assoc
18	1026	97.9	203	22	AAU02636	Synaptosomal-assoc
19	1022	97.5	202	22	AAU00265	Synaptosomal-assoc
20	1017	97.0	201	22	AAU02637	Synaptosomal-assoc
21	1012	96.6	200	22	AAU00264	Synaptosomal-assoc
22	1009	96.3	198	22	AAU00255	Synaptosomal-assoc
23	1007	96.1	199	22	AAU00263	Synaptosomal-assoc
24	1004	95.8	206	22	AAU00252	Synaptosomal-assoc
25	625.5	59.7	212	22	ABB6447	Synaptosomal-assoc
26	613.5	58.5	211	22	ABG02947	Synaptosomal-assoc
27	613.5	58.5	211	22	ABG00251	Synaptosomal-assoc
28	609.5	58.2	213	21	AAU02637	Synaptosomal-assoc
29	587	56.0	116	23	AAO15165	Synaptosomal-assoc
30	581	55.4	116	23	AAO15166	Synaptosomal-assoc
31	451	43.0	106	21	AAO03825	Synaptosomal-assoc
32	451	43.0	106	21	AAO03826	Synaptosomal-assoc
33	403	38.5	86	22	AAO15584	Synaptosomal-assoc
34	391.5	37.4	82	22	AAO15581	Synaptosomal-assoc
35	361.5	34.5	129	21	AAO53705	Synaptosomal-assoc
36	353	33.7	70	17	AAO68823	Synaptosomal-assoc
37	310	29.6	64	21	AAO00764	Synaptosomal-assoc
38	253	24.1	513	21	AAO32996	Synaptosomal-assoc
39	253	24.1	546	21	AAO32995	Synaptosomal-assoc
40	253	24.1	714	21	AAO32994	Synaptosomal-assoc
41	244	23.3	49	22	AAO57386	Synaptosomal-assoc
42	230	21.9	247	21	AAO09027	Synaptosomal-assoc
43	230	21.9	247	21	AAO33785	Synaptosomal-assoc
44	230	21.9	247	21	AAO39336	Synaptosomal-assoc
45	230	21.9	270	21	AAO23784	Synaptosomal-assoc

#### ALIGNMENTS

RESULT 1	
AAW30103	
AAW30103 standard; peptide: 206 AA.	
AC	AAW30103;
DT	06-APR-1998 (first entry)
XX	Synaptosomal associated protein.
DE	
XX	Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;
KW	excitation-secretory uncoupling peptide; catecholamine secretion;
KW	bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;
KW	synaptosomal associated protein; SNAP-25.
OS	Homo sapiens.
XX	
PN	W09734620-A1.
XX	
PD	25-SEP-1997.
XX	
PF	18-MAR-1997; 97WO-US04393.
XX	
PR	18-MAR-1996; 96US-0013599.
XX	
PA	(REGC ) UNIV CALIFORNIA.
XX	
PI	Montal M;
XX	
DR	WPI: 1997-479986/44.
XX	
PT	Excitation-secretory uncoupling peptide(s) for inhibiting
	neurotransmitter release - used particularly for treating muscle

PT spasticity, and for delivering drugs specifically to neural cells  
XX  
PS Disclosure: Page 27-28: 61pp; English.  
XX  
XX This sequence represents the human 25 kD synaptosomal associated protein  
CC (SNAP-25), which is an inhibitory agent of the invention. The agents of  
CC the invention inhibit secretion of neurotransmitter from neuronal cells  
CC and is an excitation-secretory uncoupling peptide (I) of at least 20  
CC amino acids (aa) all of which correspond substantially to any one of  
CC AAM30097-W30102, or more generally any (I) that inhibits 50% of  
CC catecholamine secretion from bovine chromaffin cells at a concentration  
CC of 10 microm, especially 0.25 microm, or less. (I) are used, as a  
CC replacement for Clostridium toxin, to inhibit release of  
CC neurotransmitters from synaptic vesicles, specifically for reducing  
CC muscle spasticity. Also (I) may be labelled to allow in vivo imaging of  
CC intracellular distribution of (I). Compounds for delivering the drug to  
CC neural cells provide targeted drug delivery, e.g. of substance P to  
CC brain tumours for induction of apoptosis. Unlike the neurotoxins, (I) are  
CC not toxic or immunogenic and are more readily available. Their  
CC therapeutic effect lasts for several days or weeks, so lower doses or  
CC less frequent treatments are required.  
XX  
SO Sequence 206 AA:  
Query Match 100.0%; Score 1048; DB 18; Length 206;  
Best Local Similarity 100.0%; Pred. No. 6,7e-91;  
Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MAEDADMRELEEMORRADOLADESLESTRMLOLVESKDGAGIRTVMLDDEGEOLERI 60  
DB 1 MAEDADMRELEEMORRADOLADESLESTRMLOLVESKDGAGIRTVMLDDEGEOLERI 60  
QY 61 EEGMDQINKDKMEAEKNTLDGKFCGLCVPCPNKLSDDAYKKAGNNDGVVASQPARV 120  
DB 61 EEGMDQINKDKMEAEKNTLDGKFCGLCVPCPNKLSDDAYKKAGNNDGVVASQPARV 120  
QY 121 VDREQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
DB 121 VDREQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
QY 181 IMEKADSNKTRIDEANORATKMLGSG 206  
DB 181 IMEKADSNKTRIDEANORATKMLGSG 206  
RESULT 2  
AAM79198  
ID AAM79198 standard; Protein: 206 AA.  
XX  
AC AAM79198;  
XX  
DT 25-NOV-1998 (first entry)  
XX  
DE Mouse SNAP-25 polypeptide.  
XX  
XX Hrs-2 polypeptide: ATP-preferring nucleotidase; SNAP-25: vesicle docking;  
XX calcium-regulated secretion; secretory vesicle; secretory process; brain;  
XX neurotransmitter release; presynaptic membrane; CNS disorder; depression;  
XX Parkinson's disease; endocrine system; hormonal imbalance; cell division;  
XX thought disorder; schizophrenia; degenerative disorder; anaesthesia; rat;  
XX immune system; antigen processing; immunomodulator; viral processing;  
XX central nervous system; vesicular release; affective disorder; human;  
XX anti-tumour application; membrane trafficking regulation; mouse.  
OS Mus sp.  
XX  
PN WO9838210-A2.  
XX  
PD 03-SEP-1998.  
XX  
PF 26-FEB-1998: 98MO-US03789.  
XX  
PR 26-FEB-1997: 97US-0039159.

XX  
PA (STRD ) UNIV IELAND STANFORD JUNIOR.  
XX  
PI Bean AJ, Scheller RH;  
XX  
XX WPI: 1998-481140/41.  
DR N-PDB: AAV57358.  
XX  
PT New isolated Hrs-2 nucleotidase - used in assays to identify  
PT compounds capable of modulating calcium-regulatory secretion of  
PT secretory vesicles, such as in neurotransmitter release  
XX  
PS Claim 16; Pages 42-44; 55pp; English.  
XX  
XX This represents a mouse SNAP-25 polypeptide, a component of the protein  
CC SNAP-25, which is an inhibitory agent of the invention. The agents of  
CC the invention inhibit secretion of neurotransmitter from neuronal cells  
CC and is an excitation-secretory uncoupling peptide (I) of at least 20  
CC amino acids (aa) all of which correspond substantially to any one of  
CC AAM30097-W30102, or more generally any (I) that inhibits 50% of  
CC catecholamine secretion from bovine chromaffin cells at a concentration  
CC of 10 microm, especially 0.25 microm, or less. (I) are used, as a  
CC replacement for Clostridium toxin, to inhibit release of  
CC neurotransmitters from synaptic vesicles, specifically for reducing  
CC muscle spasticity. Also (I) may be labelled to allow in vivo imaging of  
CC intracellular distribution of (I). Compounds for delivering the drug to  
CC neural cells provide targeted drug delivery, e.g. of substance P to  
CC brain tumours for induction of apoptosis. Unlike the neurotoxins, (I) are  
CC not toxic or immunogenic and are more readily available. Their  
CC therapeutic effect lasts for several days or weeks, so lower doses or  
CC less frequent treatments are required.  
XX  
SO Sequence 206 AA:  
Query Match 100.0%; Score 1048; DB 19; Length 206;  
Best Local Similarity 100.0%; Pred. No. 6,7e-91;  
Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MAEDADMRELEEMORRADOLADESLESTRMLOLVESKDGAGIRTVMLDDEGEOLERI 60  
DB 1 MAEDADMRELEEMORRADOLADESLESTRMLOLVESKDGAGIRTVMLDDEGEOLERI 60  
QY 61 EEGMDQINKDKMEAEKNTLDGKFCGLCVPCPNKLSDDAYKKAGNNDGVVASQPARV 120  
DB 61 EEGMDQINKDKMEAEKNTLDGKFCGLCVPCPNKLSDDAYKKAGNNDGVVASQPARV 120  
QY 121 VDREQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
DB 121 VDREQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
QY 181 IMEKADSNKTRIDEANORATKMLGSG 206  
DB 181 IMEKADSNKTRIDEANORATKMLGSG 206  
RESULT 3  
AAM43426  
ID AAM43426 standard; Protein: 206 AA.  
XX  
AC AAM43426;  
XX  
DT 27-APR-1998 (first entry)  
XX  
DE Mouse synaptosomal-associated protein-25.  
XX  
PF Binding domain: mouse; syntaxin; synaptosomal-associated protein; CNS;  
XX neurotransmitter; presynaptic membrane; central nervous system; tumour;  
KM

GenCore version 5.1.3  
Copyright (c) 1993 - 2002 Compugen Ltd.

OM protein - protein search, using sw model

Run on: November 19, 2002, 17:32:44 : Search time 3.35135 Seconds  
(without alignments)  
318.082 Million cell updates/sec

Title: PCT-US02-27145-1

Perfect score: 39

Sequence: 1 EAAQRAIK 8

Scoring table:

BLOSUM62  
Gap 10.0, Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

A\_Geneseq\_101002:\*

1: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1980.DAT.\*  
2: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1981.DAT.\*  
3: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1982.DAT.\*  
4: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1983.DAT.\*  
5: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1984.DAT.\*  
6: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1985.DAT.\*  
7: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1986.DAT.\*  
8: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1987.DAT.\*  
9: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1988.DAT.\*  
10: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1989.DAT.\*  
11: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1990.DAT.\*  
12: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1991.DAT.\*  
13: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1992.DAT.\*  
14: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1993.DAT.\*  
15: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1994.DAT.\*  
16: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1995.DAT.\*  
17: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1996.DAT.\*  
18: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1997.DAT.\*  
19: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1998.DAT.\*  
20: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1999.DAT.\*  
21: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2000.DAT.\*  
22: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2001.DAT.\*  
23: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2002.DAT.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	39	100.0	12	20	AAV44061 Human SNAP25 (amin
2	39	100.0	15	20	AAV44058 Human SNAP25 (amin
3	39	100.0	16	20	AAV44069 Human SNAP25 (amin
4	39	100.0	17	20	AAV44021 Amino acids 187-20
5	39	100.0	17	20	AAV44039 Human SNAP25 (amin
6	39	100.0	17	20	AAV44044 Human SNAP25 (amin
7	39	100.0	17	20	AAV44048 Human SNAP25 (amin
8	39	100.0	17	20	AAV44055 Human SNAP25 (amin
9	39	100.0	17	20	AAV44057 Human SNAP25 (amin
10	39	100.0	17	20	AAV44070 Human SNAP25 (amin

11	39	100.0	17	23	ABG69065
12	39	100.0	19	22	AAV5586
13	39	100.0	20	18	AAV30100
14	39	100.0	26	18	AAV30099
15	39	100.0	37	18	AAV30097
16	39	100.0	70	17	AAV86823
17	39	100.0	85	29	AAV5584
18	39	100.0	116	23	AAV51565
19	39	100.0	206	18	AAV30103
20	39	100.0	206	19	AAV79198
21	39	100.0	206	19	AAV43426
22	39	100.0	206	22	AAV00246
23	39	100.0	206	22	AAV00252
24	39	100.0	206	22	AAV00253
25	39	100.0	206	22	AAV02638
26	39	100.0	206	22	AAV02640
27	39	100.0	206	22	AAV02640
28	39	92.3	12	20	AAV44037
29	39	92.3	13	20	AAV44036
30	39	92.3	16	20	AAV44027
31	39	92.3	16	20	AAV44071
32	39	92.3	16	20	AAV44072
33	39	92.3	16	20	AAV44073
34	39	92.3	16	20	AAV44074
35	39	92.3	17	20	AAV44022
36	39	92.3	17	20	AAV44023
37	39	92.3	17	20	AAV44024
38	39	92.3	17	20	AAV44038
39	39	92.3	17	20	AAV44041
40	39	92.3	17	20	AAV44063
41	39	92.3	24	23	AAV5162
42	39	89.7	17	20	AAV44047
43	39	89.7	17	20	AAV44050
44	39	89.7	17	20	AAV44052
45	39	89.7	17	20	AAV44059

#### ALIGNMENTS

RESULT 1  
ID AAV44061 standard; peptide: 12 AA.  
AC AAV44061;  
DT 18-JAN-2000 (first entry)  
XX  
DE Human SNAP25 (amino acids 187-203) analogue #40.  
XX  
KW Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;  
KW Fluorescamine; detection; human; synaptosomal protein; SNAP25;  
KW hydrolysis; amino group.  
XX  
OS Homo sapiens.  
OS Synthetic.  
OS  
PN US5965699 A1  
XX  
PD 12-OCT-1999.  
XX  
PF 06-NOV-1996; 96US-0743894.  
XX  
PR 06-NOV-1996; 96US-0743894.  
XX  
PA (US5965699 A1)  
XX  
PI Bostian KA, Schmidt JT;  
XX WPI: 1999-57939/49.  
XX  
PT Quantitation of type A botulinum toxin -

PS Disclosure; Column 9-10; 28pp; English.

CC The invention relates to an enzymatic assay for the quantitation of  
CC type A botulinum toxin, by determining the proteolytic activity of  
CC botulinum neurotoxin type A using fluorescamine detection. Botulinum  
CC toxin A has been shown to cleave the synaptosomal neurotransmitter  
CC peptide SNAP25 between residues 197-198. The method comprises adding  
CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,  
CC amino acids 187-203 of human SNAP25) to a sample containing the  
CC botulinum toxin A so that hydrolysis of the peptide is initiated, then  
CC stopping hydrolysis of the peptide at different time points; and  
CC measuring the amount of hydrolysis at each time point by combining with a  
CC label capable of detecting free amino groups resulting from the  
CC hydrolysis. The amount of botulinum toxin A present in the sample is  
CC determined by comparing measurements with the amount of label produced  
CC from a known concentration of toxin measured under similar conditions.  
CC The method is useful for the quantitation of type A botulinum toxin.

XX Sequence 12 AA:

SO

Query Match 100.0%; Score 39; DB 20; Length 12;  
Best Local Similarity 100.0%; Pred. No. 0.066;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 EANORATK 8  
| | | | | | | |  
Db 3 EANORATK 10

RESULT 2

AAAY44058  
ID AAY44058 standard; peptide; 15 AA.

XX AAY44058;

XX 18-JAN-2000 (first entry)

DE Human SNAP25 (amino acids 187-203) analogue [1-15].

XX Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;  
KM fluorescamine; detection; human; synaptosomal protein; SNAP25;  
KM hydrolysis; amino group.

XX Homo sapiens.

XX US5965699-A.

XX 12-OCT-1999.

XX 06-NOV-1996; 96US-0743894.

XX 06-NOV-1996; 96US-0743894.

XX (USSA ) US SEC OF ARMY.

PI Bostian KA, Schmidt JI;  
XX WPI; 1999-579939/49.

XX Quantitation of type A botulinum toxin -

PT Disclosure; Column 9; 28pp; English.

XX The invention relates to an enzymatic assay for the quantitation of  
CC type A botulinum toxin, by determining the proteolytic activity of  
CC botulinum neurotoxin type A using fluorescamine detection. Botulinum  
CC toxin A has been shown to cleave the synaptosomal neurotransmitter  
CC peptide SNAP25 between residues 197-198. The method comprises adding  
CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,  
CC amino acids 187-203 of human SNAP25) to a sample containing the  
CC botulinum toxin A so that hydrolysis of the peptide is initiated, then  
CC stopping hydrolysis of the peptide at different time points; and  
CC measuring the amount of hydrolysis at each time point by combining with a

CC label capable of detecting free amino groups resulting from the  
CC hydrolysis. The amount of botulinum toxin A present in the sample is  
CC determined by comparing measurements with the amount of label produced  
CC from a known concentration of toxin measured under similar conditions.  
CC The method is useful for the quantitation of type A botulinum toxin.

XX Sequence 15 AA:

SO

Query Match 100.0%; Score 39; DB 20; Length 15;  
Best Local Similarity 100.0%; Pred. No. 0.084;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 EANORATK 8  
| | | | | | | |  
Db 8 EANORATK 15

RESULT 3

AAAY44069  
ID AAY44069 standard; peptide; 16 AA.

XX AAY44069;

XX 18-JAN-2000 (first entry)

DE Human SNAP25 (amino acids 187-203) analogue [1-16].

XX Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;  
KM fluorescamine; detection; human; synaptosomal protein; SNAP25;  
KM hydrolysis; amino group.

XX Homo sapiens.

XX US5965699-A.

XX 12-OCT-1999.

XX 06-NOV-1996; 96US-0743894.

XX 06-NOV-1996; 96US-0743894.

XX (USSA ) US SEC OF ARMY.

PI Bostian KA, Schmidt JI;  
XX WPI; 1999-579939/49.

XX Quantitation of type A botulinum toxin -

PT Disclosure; Column 13-14; 28pp; English.

XX The invention relates to an enzymatic assay for the quantitation of  
CC type A botulinum toxin, by determining the proteolytic activity of  
CC botulinum neurotoxin type A using fluorescamine detection. Botulinum  
CC toxin A has been shown to cleave the synaptosomal neurotransmitter  
CC peptide SNAP25 between residues 197-198. The method comprises adding  
CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,  
CC amino acids 187-203 of human SNAP25) to a sample containing the  
CC botulinum toxin A so that hydrolysis of the peptide is initiated, then  
CC stopping hydrolysis of the peptide at different time points; and  
CC measuring the amount of hydrolysis at each time point by combining with a  
CC label capable of detecting free amino groups resulting from the  
CC hydrolysis. The amount of botulinum toxin A present in the sample is  
CC determined by comparing measurements with the amount of label produced  
CC from a known concentration of toxin measured under similar conditions.  
CC The method is useful for the quantitation of type A botulinum toxin.

XX Sequence 16 AA:

SO

Query Match 100.0%; Score 39; DB 20; Length 16;  
Best Local Similarity 100.0%; Pred. No. 0.09;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

QY      1 EANORATK 8
XX      |||||
Db      8 EANORATK 15

RESULT 4
ID      AAY44021
XX      AAY44021 standard; peptide: 17 AA.
AC      AAY44021;
XX
XX      18-JAN-2000 (first entry)
DT
XX
XX      Amino acids 187-203 of human SNAP25.
DE
XX
XX      Enzymatic assay; quantitation: type A botulinum neurotoxin; proteolysis;
KM      fluorescamine; detection: human; synaptosomal protein; SNAP25;
KW      hydrolysis; amino group.
XX
XX      Homo sapiens.
OS
XX      US5965699-A.
PN
XX      12-OCT-1999.
PD
XX      06-NOV-1996; 96US-0743894.
PF
XX      06-NOV-1996; 96US-0743894.
PR
XX      06-NOV-1996; 96US-0743894.
XX      (USSA ) US SEC OF ARMY.
PA
XX      Bostian KA, Schmidt JJ;
PI
XX      WPI: 1999-579939/49.
XX
XX      Quantitation of type A botulinum toxin -
PT
XX      Claim 1; Column 4; 28pp; English.
PS
XX
XX      The invention relates to an enzymatic assay for the quantitation of
CC      type A botulinum toxin, by determining the proteolytic activity of
CC      botulinum neurotoxin type A using fluorescamine detection. The method
CC      comprises adding an analogue (e.g. AAY44022-Y44076) of this peptide
CC      (which represents amino acids 187-203 of the human synaptosomal protein
CC      SNAP25) to a sample containing the botulinum toxin A so that hydrolysis
CC      of the peptide is initiated, then stopping hydrolysis of the peptide at
CC      different time points; and measuring the amount of hydrolysis at each
CC      time point by combining with a label capable of detecting free amino
CC      groups resulting from the hydrolysis. The amount of botulinum toxin A
CC      present in the sample is determined by comparing measurements with the
CC      amount of label produced from a known concentration of toxin measured
CC      under similar conditions. The method is useful for the quantitation of
CC      type A botulinum toxin.
XX
XX      Sequence 17 AA:
SQ
XX
XX      Query Match 100.0%; Score 39; DB 20; Length 17;
XX      Best Local Similarity 100.0%; Pred. No. 0.097;
XX      Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 EANORATK 8
XX      |||||
Db      8 EANORATK 15

RESULT 5
ID      AAY44039
XX      AAY44039 standard; peptide: 17 AA.
AC
XX      AAY44039;
XX
XX      18-JAN-2000 (first entry)
DT
XX

```

```

DE      Human SNAP25 (amino acids 187-203) analogue #18.
XX
XX      Enzymatic assay; quantitation: type A botulinum neurotoxin; proteolysis;
KM      fluorescamine; detection: human; synaptosomal protein; SNAP25;
KW      hydrolysis; amino group.
XX
XX      Homo sapiens.
OS
XX      Synthetic.
XX      Key Modified-site 4 Location/Qualifiers
XX      FT /label= Abu
XX
XX      US5965699-A.
PN
XX      12-OCT-1999.
PD
XX      06-NOV-1996; 96US-0743894.
PF
XX      06-NOV-1996; 96US-0743894.
PR
XX      06-NOV-1996; 96US-0743894.
XX      (USSA ) US SEC OF ARMY.
PA
XX      Bostian KA, Schmidt JJ;
PI
XX      WPI: 1999-579939/49.
XX
XX      Quantitation of type A botulinum toxin -
PT
XX      Disclosure; Column 7-8; 28pp; English.
PS
XX
XX      The invention relates to an enzymatic assay for the quantitation of
CC      type A botulinum toxin, by determining the proteolytic activity of
CC      botulinum neurotoxin type A using fluorescamine detection. Botulinum
CC      toxin A has been shown to cleave the synaptosomal neurotransmitter
CC      peptide SNAP25 between residues 197-198. The method comprises adding
CC      an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,
CC      amino acids 187-203 of human SNAP25) to a sample containing the
CC      botulinum toxin A so that hydrolysis of the peptide is initiated, then
CC      stopping hydrolysis of the peptide at different time points; and
CC      measuring the amount of hydrolysis at each time point by combining with a
CC      label capable of detecting free amino groups resulting from the
CC      hydrolysis. The amount of botulinum toxin A present in the sample is
CC      determined by comparing measurements with the amount of label produced
CC      from a known concentration of toxin measured under similar conditions.
CC      The method is useful for the quantitation of type A botulinum toxin.
XX
XX      Sequence 17 AA:
SQ
XX
XX      Query Match 100.0%; Score 39; DB 20; Length 17;
XX      Best Local Similarity 100.0%; Pred. No. 0.097;
XX      Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 EANORATK 8
XX      |||||
Db      8 EANORATK 15

RESULT 6
ID      AAY44044
XX      AAY44044 standard; peptide: 17 AA.
AC
XX      AAY44044;
XX
XX      18-JAN-2000 (first entry)
DT
XX
XX      Human SNAP25 (amino acids 187-203) analogue M16X.
DE
XX
XX      Enzymatic assay; quantitation: type A botulinum neurotoxin; proteolysis;
KM      fluorescamine; detection: human; synaptosomal protein; SNAP25;
KW      hydrolysis; amino group.
XX
XX      Homo sapiens.
OS

```

OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 16 /label= Nle  
XX  
XX US5965699-A.  
XX  
PD 12-OCT-1999.  
XX  
PF 06-NOV-1996; 96US-0743894.  
XX  
XX 06-NOV-1996; 96US-0743894.  
XX  
XX (USSA ) US SEC OF ARMY.  
XX  
XX Bostian KA, Schmidt JJ;  
PI  
PI WPI; 1999-579939/49.  
XX  
XX  
XX Quantitation of type A botulinum toxin -  
PT  
PS Disclosure; Column 7-8; 28pp; English.  
XX  
XX The invention relates to an enzymatic assay for the quantitation of  
CC type A botulinum toxin, by determining the proteolytic activity of  
CC botulinum neurotoxin type A using fluorescamine detection. Botulinum  
CC toxin A has been shown to cleave the synaptosomal neurotransmitter  
CC peptide SNAP25 between residues 197-198. The method comprises adding  
CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,  
CC amino acids 187-203 of human SNAP25) to a sample containing the  
CC botulinum toxin A so that hydrolysis of the peptide is initiated, then  
CC stopping hydrolysis of the peptide at different time points; and  
CC measuring the amount of hydrolysis at each time point by combining with a  
CC label capable of detecting free amino groups resulting from the  
CC hydrolysis. The amount of botulinum toxin A present in the sample is  
CC determined by comparing measurements with the amount of label produced  
CC from a known concentration of toxin measured under similar conditions.  
CC The method is useful for the quantitation of type A botulinum toxin.  
XX  
SQ Sequence 17 AA:  
  
Query Match 100.0%; Score 39; DB 20; Length 17;  
Best Local Similarity 100.0%; Pred. No. 0.097;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
OY 1 EANORATK 8  
DB 8 EANORATK 15.  
  
RESULT 7  
AAY44048  
ID AAY44048 standard; peptide; 17 AA.  
XX  
XX AAY44048;  
AC  
XX  
DT 18-JAN-2000 (first entry)  
XX  
DE Human SNAP25 (amino acids 187-203) analogue M16A.  
XX  
XX Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;  
KW fluorescamine; detection; human; synaptosomal protein; SNAP25;  
KW hydrolysis; amino group.  
XX  
XX Homo sapiens.  
OS Synthetic.  
XX  
XX US5965699-A.  
PN  
XX  
PD 12-OCT-1999.  
XX  
PF 06-NOV-1996; 96US-0743894.

XX  
XX 06-NOV-1996; 96US-0743894.  
XX  
XX (USSA ) US SEC OF ARMY.  
PA  
XX  
XX Bostian KA, Schmidt JJ;  
PI  
PI WPI; 1999-579939/49.  
XX  
XX  
XX Quantitation of type A botulinum toxin -  
PT  
PS Disclosure; Column 9; 28pp; English.  
XX  
XX The invention relates to an enzymatic assay for the quantitation of  
CC type A botulinum toxin, by determining the proteolytic activity of  
CC botulinum neurotoxin type A using fluorescamine detection. Botulinum  
CC toxin A has been shown to cleave the synaptosomal neurotransmitter  
CC peptide SNAP25 between residues 197-198. The method comprises adding  
CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,  
CC amino acids 187-203 of human SNAP25) to a sample containing the  
CC botulinum toxin A so that hydrolysis of the peptide is initiated, then  
CC stopping hydrolysis of the peptide at different time points; and  
CC measuring the amount of hydrolysis at each time point by combining with a  
CC label capable of detecting free amino groups resulting from the  
CC hydrolysis. The amount of botulinum toxin A present in the sample is  
CC determined by comparing measurements with the amount of label produced  
CC from a known concentration of toxin measured under similar conditions.  
CC The method is useful for the quantitation of type A botulinum toxin.  
XX  
SQ Sequence 17 AA:  
  
Query Match 100.0%; Score 39; DB 20; Length 17;  
Best Local Similarity 100.0%; Pred. No. 0.097;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
OY 1 EANORATK 8  
DB 8 EANORATK 15  
  
RESULT 8  
AAY44056  
ID AAY44056 standard; peptide; 17 AA.  
XX  
XX AAY44056;  
AC  
XX  
DT 18-JAN-2000 (first entry)  
XX  
DE Human SNAP25 (amino acids 187-203) analogue #35.  
XX  
XX Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;  
KW fluorescamine; detection; human; synaptosomal protein; SNAP25;  
KW hydrolysis; amino group.  
XX  
XX Homo sapiens.  
OS Synthetic.  
XX  
XX US5965699-A.  
PN  
XX  
PD 12-OCT-1999.  
XX  
PF 06-NOV-1996; 96US-0743894.  
XX  
XX 06-NOV-1996; 96US-0743894.  
XX  
XX (USSA ) US SEC OF ARMY.  
XX  
XX Bostian KA, Schmidt JJ;  
PI  
PI WPI; 1999-579939/49.  
XX  
XX Quantitation of type A botulinum toxin -  
PT



PS Disclosure; Column 9; 28pp; English.

XX The invention relates to an enzymatic assay for the quantitation of

CC type A botulinum toxin, by determining the proteolytic activity of

CC botulinum neurotoxin type A using fluorescamine detection. Botulinum

CC toxin A has been shown to cleave the synaptosomal neurotransmitter

CC peptide SNAP25 between residues 197-198. The method comprises adding

CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,

CC amino acids 187-203 of human SNAP25) to a sample containing the

CC botulinum toxin A so that hydrolysis of the peptide is initiated, then

CC stopping hydrolysis of the peptide at different time points; and

CC measuring the amount of hydrolysis at each time point by combining with a

CC label capable of detecting free amino groups resulting from the

CC hydrolysis. The amount of botulinum toxin A present in the sample is

CC determined by comparing measurements with the amount of label produced

CC from a known concentration of toxin measured under similar conditions.

CC The method is useful for the quantitation of type A botulinum toxin.

CC

XX Sequence 17 AA:

SQ

Query Match 100.0%; Score 39; DB 20; Length 17;

Best Local Similarity 100.0%; Pred. No. 0.097;

Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 EANORATK 8

IIIIIIII

DB 8 EANORATK 15

XX

RESULT 9

AAY44057

ID AAY44057 standard; peptide; 17 AA.

XX

AC AAY44057:

XX

DT 18-JAN-2000 (first entry)

XX

DE Human SNAP25 (amino acids 187-203) analogue #36.

XX

KW Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;

KW fluorescamine; detection; human; synaptosomal protein; SNAP25;

KW hydrolysis; amino group.

XX

OS Homo sapiens.

OS Synthetic.

XX

PN US5965699-A.

XX

PD 12-OCT-1999.

XX

PF 06-NOV-1996; 96US-0743894.

XX

PR 06-NOV-1996; 96US-0743894.

XX

PS (USSA ) US SEC OF ARMY.

XX

PI Bostian KA, Schmidt JT;

XX

DR WPI: 1999-579939/49.

XX

PT Quantitation of type A botulinum toxin -

XX

PS Disclosure; Column 9; 28pp; English.

XX

CC The invention relates to an enzymatic assay for the quantitation of

CC type A botulinum toxin, by determining the proteolytic activity of

CC botulinum neurotoxin type A using fluorescamine detection. Botulinum

CC toxin A has been shown to cleave the synaptosomal neurotransmitter

CC peptide SNAP25 between residues 197-198. The method comprises adding

CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,

CC amino acids 187-203 of human SNAP25) to a sample containing the

CC botulinum toxin A so that hydrolysis of the peptide is initiated, then

CC stopping hydrolysis of the peptide at different time points; and

CC measuring the amount of hydrolysis at each time point by combining with a

CC label capable of detecting free amino groups resulting from the

CC hydrolysis. The amount of botulinum toxin A present in the sample is

CC determined by comparing measurements with the amount of label produced

CC from a known concentration of toxin measured under similar conditions.

CC The method is useful for the quantitation of type A botulinum toxin.

CC

XX Sequence 17 AA:

SQ

Query Match 100.0%; Score 39; DB 20; Length 17;

Best Local Similarity 100.0%; Pred. No. 0.097;

Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 EANORATK 8

IIIIIIII

DB 8 EANORATK 15

XX

RESULT 10

AAY44070

ID AAY44070 standard; peptide; 17 AA.

XX

AC AAY44070:

XX

DT 18-JAN-2000 (first entry)

XX

DE Human SNAP25 (amino acids 187-203) analogue D7N.

XX

KW Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;

KW fluorescamine; detection; human; synaptosomal protein; SNAP25;

KW hydrolysis; amino group.

XX

OS Homo sapiens.

OS Synthetic.

XX

PN US5965699-A.

XX

PD 12-OCT-1999.

XX

PF 06-NOV-1996; 96US-0743894.

XX

PR 06-NOV-1996; 96US-0743894.

XX

PS (USSA ) US SEC OF ARMY.

XX

PI Bostian KA, Schmidt JT;

XX

DR WPI: 1999-579939/49.

XX

PT Quantitation of type A botulinum toxin -

XX

PS Disclosure; Column 15; 28pp; English.

XX

CC The invention relates to an enzymatic assay for the quantitation of

CC type A botulinum toxin, by determining the proteolytic activity of

CC botulinum neurotoxin type A using fluorescamine detection. Botulinum

CC toxin A has been shown to cleave the synaptosomal neurotransmitter

CC peptide SNAP25 between residues 197-198. The method comprises adding

CC an analogue (e.g. AAY44022-Y44076) of the SNAP25 peptide (AAY44021,

CC amino acids 187-203 of human SNAP25) to a sample containing the

CC botulinum toxin A so that hydrolysis of the peptide is initiated, then

CC stopping hydrolysis of the peptide at different time points; and

CC measuring the amount of hydrolysis at each time point by combining with a

CC label capable of detecting free amino groups resulting from the

CC hydrolysis. The amount of botulinum toxin A present in the sample is

CC determined by comparing measurements with the amount of label produced

CC from a known concentration of toxin measured under similar conditions.

CC The method is useful for the quantitation of type A botulinum toxin.

CC

XX Sequence 17 AA:

SQ

Query Match 100.0%; Score 39; DB 20; Length 17;

Best Local Similarity 100.0%; Pred. No. 0.097;



KW synaptosomal associated protein; SNAP-25.  
 XX Homo sapiens.  
 XX WO9734620-A1.  
 XX 25-SEP-1997.  
 XX 18-MAR-1997; 97WO-US04393.  
 XX 18-MAR-1996; 96US-0013599.  
 XX (REGC ) UNIV CALIFORNIA.  
 XX Montal M;  
 XX WPI; 1997-479986/44.  
 XX Excitation-secretory uncoupling peptide(s) for inhibiting  
 PT neuro:transmitter release - used particularly for treating muscle  
 PT spasticity, and for delivering drugs specifically to neural cells  
 XX  
 PS Claim 14; Page 32; 61pp; English.  
 XX This sequence corresponds to residues 187-206 of the human 25 kD  
 CC synaptosomal associated protein (SNAP-25), and is an inhibitory agent of  
 CC the invention. The agents of the invention inhibit secretion of  
 CC neurotransmitter from neuronal cells and is an excitation-secretory  
 CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which  
 CC correspond substantially to any one of AAM30097-W30102, or more  
 CC generally any (I) that inhibits 50% of catecholamine secretion from  
 CC bovine chromaffin cells at a concentration of 10 microm, especially 0.25  
 CC microm, or less. (I) are used, as a replacement for Clostridium toxin, to  
 CC inhibit release of neurotransmitters from synaptic vesicles, specifically  
 CC for reducing muscle spasticity. Also (I) may be labelled to allow in  
 CC vivo imaging of intracellular distribution of (I). Compounds for  
 CC delivering the drug to neural cells provide targeted drug delivery, e.g.  
 CC of substance P to brain tumours for induction of apoptosis. Unlike the  
 CC neurotoxins, (I) are not toxic or immunogenic and are more readily  
 CC available. Their therapeutic effect lasts for several days or weeks, so  
 CC lower doses or less frequent treatments are required.  
 XX  
 SQ Sequence 20 AA;  
 Query Match 100.0%; Score 39; DB 18; Length 20;  
 Best Local Similarity 100.0%; Pred. No. 0.12;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 OY 1 EANORATK 8  
 Db 8 EANORATK 15  
 RESULT 14  
 AAM30099  
 ID AAM30099 standard; peptide: 26 AA.  
 XX AAM30099;  
 XX 06-APR-1998 (first entry)  
 XX Neurotransmitter secretion inhibitor #3.  
 XX Neurotransmitter secretion inhibitor; neuronal cell; synaptic vesicle;  
 KW excitation-secretory uncoupling peptide; catecholamine secretion;  
 KW bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
 KW synaptosomal associated protein; SNAP-25.  
 XX Homo sapiens.  
 XX WO9734620-A1.  
 XX 25-SEP-1997.

XX 18-MAR-1997; 97WO-US04393.  
 XX 18-MAR-1996; 96US-0013599.  
 XX (REGC ) UNIV CALIFORNIA.  
 XX Montal M;  
 XX WPI; 1997-479986/44.  
 XX Excitation-secretory uncoupling peptide(s) for inhibiting  
 PT neuro:transmitter release - used particularly for treating muscle  
 PT spasticity, and for delivering drugs specifically to neural cells  
 XX  
 PS Claim 13; Page 31; 61pp; English.  
 XX This sequence corresponds to residues 181-206 of the human 25 kD  
 CC synaptosomal associated protein (SNAP-25), and is an inhibitory agent of  
 CC the invention. The agents of the invention inhibit secretion of  
 CC neurotransmitter from neuronal cells and is an excitation-secretory  
 CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which  
 CC correspond substantially to any one of AAM30097-W30102, or more  
 CC generally any (I) that inhibits 50% of catecholamine secretion from  
 CC bovine chromaffin cells at a concentration of 10 microm, especially 0.25  
 CC microm, or less. (I) are used, as a replacement for Clostridium toxin, to  
 CC inhibit release of neurotransmitters from synaptic vesicles, specifically  
 CC for reducing muscle spasticity. Also (I) may be labelled to allow in  
 CC vivo imaging of intracellular distribution of (I). Compounds for  
 CC delivering the drug to neural cells provide targeted drug delivery, e.g.  
 CC of substance P to brain tumours for induction of apoptosis. Unlike the  
 CC neurotoxins, (I) are not toxic or immunogenic and are more readily  
 CC available. Their therapeutic effect lasts for several days or weeks, so  
 CC lower doses or less frequent treatments are required.  
 XX  
 SQ Sequence 26 AA;  
 Query Match 100.0%; Score 39; DB 18; Length 26;  
 Best Local Similarity 100.0%; Pred. No. 0.16;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 OY 1 EANORATK 8  
 Db 14 EANORATK 21  
 RESULT 15  
 AAM30097  
 ID AAM30097 standard; peptide: 37 AA.  
 XX AAM30097;  
 XX 06-APR-1998 (first entry)  
 XX Neurotransmitter secretion inhibitor #1.  
 XX Neurotransmitter secretion inhibitor; neuronal cell; synaptic vesicle;  
 KW excitation-secretory uncoupling peptide; catecholamine secretion;  
 KW bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
 KW synaptosomal associated protein; SNAP-25.  
 XX Homo sapiens.  
 XX WO9734620-A1.  
 XX 25-SEP-1997.  
 XX 18-MAR-1997; 97WO-US04393.  
 XX 18-MAR-1996; 96US-0013599.  
 XX (REGC ) UNIV CALIFORNIA.

PI Montal M;  
XX  
DR WPI: 1997-479986/44.

XX  
PT Excitation-secretory uncoupling peptide(s) for inhibiting  
PT neuro:transmitter release - used particularly for treating muscle  
PT spasticity, and for delivering drugs specifically to neural cells  
XX

PS Claim 1: Page 30: 61pp; English.

XX  
CC This sequence corresponds to residues 170-206 of the human 25 kD  
CC synaptosomal associated protein (SNAP-25), and is a inhibitory agent of  
CC the invention. The agents of the invention inhibit secretion of  
CC neurotransmitter from neuronal cells and is an excitation-secretory  
CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which  
CC correspond substantially to any one of AAM30097-W30102, or more  
CC generally any (I) that inhibits 50% of catecholamine secretion from  
CC bovine chromaffin cells at a concentration of 10 microm, especially 0.25  
CC microm, or less. (I) are used, as a replacement for Clostridium toxin, to  
CC inhibit release of neurotransmitters from synaptic vesicles, specifically  
CC for reducing muscle spasticity. Also (I) may be labelled to allow in  
CC vivo imaging of intracellular distribution of (I). Compounds for  
CC delivering the drug to neural cells provide targeted drug delivery, e.g.  
CC of substance P to brain tumours for induction of apoptosis. Unlike the  
CC neurotoxins, (I) are not toxic or immunogenic and are more readily  
CC available. Their therapeutic effect lasts for several days or weeks, so  
CC lower doses or less frequent treatments are required.

XX  
SQ Sequence 37 AA:

Query Match 100.0%; Score 39; DB 18; Length 37;  
Best Local Similarity 100.0%; Pred. No. 0.23;

Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 EANQRA TK 8  
IIIIIIII  
DB 25 EANQRA TK 32

Search completed: November 19, 2002, 17:34:25  
Job time : 4.35135 secs

GenCore version 5.1.3  
Copyright (c) 1993 - 2002 Compugen Ltd.

OM protein - protein search, using sw model

Run on: November 19, 2002, 17:32:44 : Search time 86.2973 Seconds  
(without alignments)  
318.082 Million cell updates/sec

Title: PCT-US02-27145-2

Perfect score: 1048  
Sequence: 1 MAEDADMRNELEEMQRRAQ.....SNKTRIDEANQRATKMLGSG 206

Scoring table:

BIOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

A\_Geneseq\_101002:\*

1: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1980.DAT:\*  
2: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1981.DAT:\*  
3: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1982.DAT:\*  
4: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1983.DAT:\*  
5: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1984.DAT:\*  
6: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1985.DAT:\*  
7: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1986.DAT:\*  
8: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1987.DAT:\*  
9: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1988.DAT:\*  
10: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1989.DAT:\*  
11: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1990.DAT:\*  
12: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1991.DAT:\*  
13: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1992.DAT:\*  
14: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1993.DAT:\*  
15: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1994.DAT:\*  
16: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1995.DAT:\*  
17: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1996.DAT:\*  
18: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1997.DAT:\*  
19: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1998.DAT:\*  
20: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA1999.DAT:\*  
21: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA2000.DAT:\*  
22: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA2001.DAT:\*  
23: /SIDS2/gcgcdata/geneseq/geneseqp-emb1/AA2002.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1048	100.0	206	AAW30103	Synaptosomal assoc
2	1048	100.0	206	AAW79198	Mouse SNAP-25 poly
3	1048	100.0	206	AAW43426	Mouse synaptosomal
4	1048	100.0	206	AAU00246	Synaptosomal-assoc
5	1048	100.0	206	AAU00253	SNARE homologue, s
6	1043	99.5	206	AAU02640	Synaptosomal-assoc
7	1042	99.4	206	AAU00259	Synaptosomal-assoc
8	1042	99.4	206	AAU00260	Synaptosomal-assoc
9	1042	99.4	206	AAU00261	Synaptosomal-assoc
10	1042	99.4	206	AAU02638	Synaptosomal-assoc

11	1041	99.3	206	AAU00262	Synaptosomal-assoc
12	1039	99.1	206	AAU00257	Synaptosomal-assoc
13	1039	99.1	206	AAU00266	Synaptosomal-assoc
14	1037	99.0	206	AAU02171	Synaptosomal-assoc
15	1036	98.9	206	AAU00256	Synaptosomal-assoc
16	1036	98.9	206	AAU02639	Synaptosomal-assoc
17	1033	98.6	206	AAU00258	Synaptosomal-assoc
18	1026	97.9	203	AAU02636	Synaptosomal-assoc
19	1022	97.5	202	AAU00265	Synaptosomal-assoc
20	1017	97.0	201	AAU02637	Synaptosomal-assoc
21	1012	96.6	200	AAU00264	Synaptosomal-assoc
22	1009	96.3	198	AAU00255	Synaptosomal-assoc
23	1007	96.1	199	AAU00263	Synaptosomal-assoc
24	1004	95.8	206	AAU00252	SNARE homologue, s
25	625.5	59.7	212	ABR64447	Drosophila melanog
26	613.5	58.5	211	ABG02947	Novel human diago
27	613.5	58.5	211	AAU00251	SNARE homologue, s
28	609.5	58.2	213	AAU00251	Human prostate can
29	587	56.0	116	AAO15165	Clostridial neurot
30	581	55.4	116	AAO15166	Clostridial neurot
31	451	43.0	106	AAO38825	Human secreted pro
32	451	43.0	106	AAO38826	Human secreted pro
33	403	38.5	86	AAU15584	Human SNAP-25 N-te
34	391.5	37.4	82	AAU15581	Human colon cancer
35	361.5	34.5	129	AAU33705	SNAP-25 residues 1
36	353	33.7	70	AAU86823	Human secreted pro
37	310	29.6	64	AAU00764	Arabidopsis thalia
38	253	24.1	513	AAU32996	Arabidopsis thalia
39	253	24.1	546	AAU32995	Arabidopsis thalia
40	253	24.1	714	AAU32994	Arabidopsis thalia
41	244	23.3	49	AAU57386	Human brain expres
42	230	21.9	247	AAU09027	Arabidopsis thalia
43	230	21.9	247	AAU33785	Arabidopsis thalia
44	230	21.9	247	AAU33386	Arabidopsis thalia
45	230	21.9	270	AAU33784	Arabidopsis thalia

#### ALIGNMENTS

RESULT 1  
AAW30103  
ID AAW30103 standard; peptide: 206 AA.  
XX AC AAW30103;  
XX DT 06-APR-1998 (first entry)  
XX DE Synaptosomal associated protein.  
XX KW Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;  
KW excitation-secretory uncoupling peptide; catecholamine secretion;  
KW bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
KW synaptosomal associated protein; SNAP-25.  
XX OS Homo sapiens.  
XX PN W09734620-A1.  
XX PD 25-SEP-1997.  
XX PF 18-MAR-1997; 97WO-US04393.  
XX PR 18-MAR-1996; 96US-0013599.  
XX PA (REGC) UNIV CALIFORNIA.  
XX PI Montal M;  
XX WP1: 1997-479986/44.  
XX PT Excitation-secretory uncoupling peptide(s) for inhibiting  
PT neuro:transmitter release - used particularly for treating muscle

PT spasticity, and for delivering drugs specifically to neural cells  
XX  
XX  
PS Disclosure: Page 27-28; 61pp; English.  
XX  
XX This sequence represents the human 25 kD synaptosomal associated protein  
CC (SNAP-25), which is an inhibitory agent of the invention. The agents of  
CC the invention inhibit secretion of neurotransmitter from neuronal cells  
CC and is an excitation-secretion uncoupling peptide (I) of at least 20  
CC amino acids (aa) all of which correspond substantially to any one of  
CC AAW0097-W0102, or more generally any (I) that inhibits 50% of  
CC catecholamine secretion from bovine chromaffin cells at a concentration  
CC of 10 microm, especially 0.25 microm, or less. (I) are used, as a  
CC replacement for Clostridium toxin, to inhibit release of  
CC neurotransmitters from synaptic vesicles, specifically for reducing  
CC muscle spasticity. Also (I) may be labelled to allow in vivo imaging of  
CC intracellular distribution of (I). Compounds for delivering the drug to  
CC neural cells provide targeted drug delivery, e.g. of substance P to  
CC brain tumours for induction of apoptosis. Unlike the neurotoxins, (I) are  
CC not toxic or immunogenic and are more readily available. Their  
CC therapeutic effect lasts for several days or weeks, so lower doses or  
CC less frequent treatments are required.  
CC  
XX  
XX  
SO Sequence 206 AA:  
  
Query Match 100.0%; Score 1048; DB 18; Length 206;  
Best Local Similarity 100.0%; Pred. No. 6,7e-91;  
Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
OY 1 MADDADMRNLEEMORADOLADESLESTRMLQVLESKDACIRTLVMDDEGEQLERI 60  
DB 1 MADDADMRNLEEMORADOLADESLESTRMLQVLESKDACIRTLVMDDEGEQLERI 60  
OY 61 EEGMDQINKDKMEAEKNTLDLGFGLGVCPCNKLSKSDAYKKAMGNNODGVVASQPARV 120  
DB 61 EEGMDQINKDKMEAEKNTLDLGFGLGVCPCNKLSKSDAYKKAMGNNODGVVASQPARV 120  
OY 121 VDEREQMAISGGFIRRYTNDARENEMDENLEQVSGIIGNLRHMAIDMGNEIDTQNRIDR 180  
DB 121 VDEREQMAISGGFIRRYTNDARENEMDENLEQVSGIIGNLRHMAIDMGNEIDTQNRIDR 180  
OY 181 IMEKADSNKTRIDEANORATKMLGSG 206  
DB 181 IMEKADSNKTRIDEANORATKMLGSG 206  
  
RESULT 2  
AAW79198  
ID AAW79198 standard; Protein: 206 AA.  
XX  
XX  
AC AAW79198:  
XX  
XX  
DT 25-NOV-1998 (first entry)  
XX  
DE Mouse SNAP-25 polypeptide.  
XX  
XX Hrs-2 polypeptide: ATP-prefering nucleotidase; SNAP-25; vesicle docking;  
KM calcium-regulated secretion; secretory vesicle; secretory process; brain;  
KM neurotransmitter release; presynaptic membrane; CNS disorder; depression;  
KM Parkinson's disease; endocrine system; hormonal imbalance; cell division;  
KM thought disorder; schizophrenia; degenerative disorder; anaesthesia; rat;  
KM immune system; antigen processing; immunomodulator; viral processing;  
KM central nervous system; vesicular release; affective disorder; human;  
KM anti-tumour application; membrane trafficking regulation; mouse.  
XX  
XX  
OS Mus sp.  
XX  
XX  
PN W09838210-A2.  
XX  
XX  
PD 03-SEP-1998.  
XX  
XX  
PF 26-FEB-1998; 98WO-US03789.  
XX  
XX  
PR 26-FEB-1997; 97US-0039159.

XX  
PA (STPD ) UNIV LELAND STANFORD JUNIOR.  
XX  
XX Bean AJ, Schellier RH;  
XX  
XX WPI: 1998-481140/41.  
DR N-PSDB: AAW57558.  
XX  
XX  
PT New isolated Hrs-2 nucleotidase - used in assays to identify  
PT compounds capable of modulating calcium-regulatory secretion of  
PT secretory vesicles, such as in neurotransmitter release  
XX  
XX  
PS Claim 16; Pages 42-44; 55pp; English.  
XX  
XX This represents a mouse SNAP-25 polypeptide, a component of the protein  
CC polypeptides thought to underlie vesicle docking and fusion. The  
CC invention provides rat and human Hrs-2 polypeptides which are ATP-  
CC preferring nucleotidase that associate with SNAP-25. For identifying a  
CC compound capable of modulating calcium-regulated secretion of secretory  
CC vesicles, a SNAP-25 polypeptide can be contacted with a Hrs-2  
CC polypeptide, in the presence and absence of a test compound. The effect  
CC of the test compound on the extent of binding between the SNAP-25 and  
CC Hrs-2 polypeptides are measured and a compound is identified as effective  
CC if its measured effect on the extent of binding is above a threshold  
CC level. The products can be used for identifying drugs capable of  
CC affecting secretory processes, such as neurotransmitter release at the  
CC active zones of presynaptic membranes. Such drugs can be used for  
CC treating disorders or conditions of the central nervous system by  
CC selectively enhancing or inhibiting vesicular release in specific areas  
CC of the brain, including affective disorders (e.g. depression), disorders  
CC of thought (e.g. schizophrenia) and degenerative disorders (Parkinson's  
CC disease), as well as applications such as anaesthesia. The drugs can  
CC also be used therapeutically in other systems such as the endocrine  
CC system for treatment of hormonal imbalances, the immune system for  
CC intervention in antigen processing, secreted immunomodulators, and viral  
CC processing, as well as anti-tumour applications, such as regulation of  
CC membrane trafficking during rapid cell division.  
XX  
XX  
SO Sequence 206 AA:  
  
Query Match 100.0%; Score 1048; DB 18; Length 206;  
Best Local Similarity 100.0%; Pred. No. 6,7e-91;  
Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
OY 1 MADDADMRNLEEMORADOLADESLESTRMLQVLESKDACIRTLVMDDEGEQLERI 60  
DB 1 MADDADMRNLEEMORADOLADESLESTRMLQVLESKDACIRTLVMDDEGEQLERI 60  
OY 61 EEGMDQINKDKMEAEKNTLDLGFGLGVCPCNKLSKSDAYKKAMGNNODGVVASQPARV 120  
DB 61 EEGMDQINKDKMEAEKNTLDLGFGLGVCPCNKLSKSDAYKKAMGNNODGVVASQPARV 120  
OY 121 VDEREQMAISGGFIRRYTNDARENEMDENLEQVSGIIGNLRHMAIDMGNEIDTQNRIDR 180  
DB 121 VDEREQMAISGGFIRRYTNDARENEMDENLEQVSGIIGNLRHMAIDMGNEIDTQNRIDR 180  
OY 181 IMEKADSNKTRIDEANORATKMLGSG 206  
DB 181 IMEKADSNKTRIDEANORATKMLGSG 206  
  
RESULT 3  
AAW43426  
ID AAW43426 standard; Protein: 206 AA.  
XX  
XX  
AC AAW43426:  
XX  
XX  
DT 27-APR-1998 (first entry)  
XX  
DE Mouse synaptosomal-associated protein-25.  
XX  
XX Blinding domain; mouse; syntaxin; synaptosomal-associated protein; CNS;  
KM neurotransmitter; presynaptic membrane; central nervous system; tumour;

KW neurodegenerative disease; hormonal disorder; immunological disorder.  
 XX Mus sp.  
 OS  
 XX US5693476-A.  
 PN  
 XX 02-DEC-1997.  
 PD  
 XX 24-FEB-1995; 95US-0393985.  
 PF  
 XX 24-FEB-1995; 95US-0393985.  
 PR  
 XX 24-FEB-1995; 95US-0393985.  
 PA (STRD ) UNIV LELAND STANFORD JUNIOR.  
 PI Scheller RH;  
 DR WPI: 1998-031743/03.  
 N-PSDB: AAV01554.  
 XX  
 XX Screening assay for modulators of syntaxin binding - using peptide  
 PT comprising binding site of syntaxin, for identifying drugs useful  
 PT for treating CNS disorders, neuro-degenerative diseases, etc  
 XX  
 XX Disclosure: Column 67-72; 57pp; English.  
 PS  
 XX This amino acid sequence represents the mouse synaptosomal-associated  
 CC protein of 25 kD (SNAP-25). The invention relates to a method for  
 CC identifying a compound capable of affecting the binding of a  
 CC syntaxin-binding protein (SBP), e.g. SNAP-25, alpha-SNAP, n-secl or VAMP,  
 CC to syntaxin. The method comprises measuring the effect of the test  
 CC compound on the extent of binding between the SBP and the SBP-binding  
 CC site on syntaxin. The method can be used for identifying drugs capable  
 CC of inhibiting or stimulating neurotransmitter release at the active zones  
 CC of presynaptic membranes, which may be useful for treating CNS disorders,  
 CC affective or psychotic disorders, neurodegenerative diseases, hormonal or  
 CC immunological disorders or tumours.  
 CC  
 SQ Sequence 206 AA;  
 Query Match 100.0%; Score 1048; DB 19; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 6,7e-91;  
 Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVLEESKDAIGRTIVYALDEOGQQLERI 60  
 DB 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVLEESKDAIGRTIVYALDEOGQQLERI 60  
 QY 61 EEGMDQINKDKMEAEKKNITLDGKFCGLCYVCPCKNKLKSSDAYKKAGNNDGVAASOPARV 120  
 DB 61 EEGMDQINKDKMEAEKKNITLDGKFCGLCYVCPCKNKLKSSDAYKKAGNNDGVAASOPARV 120  
 QY 121 VEREOMATISGGFIRRYTVDARENEMDENLEOVSGITIGLRRHMLDMGNEIDTQNRQIDR 180  
 DB 121 VEREOMATISGGFIRRYTVDARENEMDENLEOVSGITIGLRRHMLDMGNEIDTQNRQIDR 180  
 QY 181 IMEKADSNTKRIDEANORATKMLGSG 206  
 DB 181 IMEKADSNTKRIDEANORATKMLGSG 206  
 RESULT 4  
 AAU000246  
 ID AAU000246 standard; Protein; 206 AA.  
 XX  
 AC AAU000246;  
 XX  
 XX 12-SEP-2001 (first entry)  
 DT  
 XX  
 DE Synaptosomal-associated protein, SNAP25.  
 XX  
 KM SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptosomal-associated protein; mutagenic; PCR primer; mouse;

KW N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 XX Mus sp.  
 OS  
 XX  
 XX Key Location/Qualifiers  
 FH Cleavage-site 180..181  
 FT Cleavage-site /note= "peptide bond susceptible to cleavage by  
 FT clostridial neurotoxin"  
 FT Cleavage-site 197..198  
 FT /note= "peptide bonds susceptible to cleavage by  
 FT clostridial neurotoxin"  
 XX  
 XX WO200118038-A2.  
 XX  
 XX 15-MAR-2001.  
 PD  
 XX 18-AUG-2000; 2000WO-GB03196.  
 PF  
 XX 20-AUG-1999; 99US-0149993.  
 PR  
 XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 PA Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
 PI WPI: 2001-226739/23.  
 DR  
 XX Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -  
 XX  
 XX Disclosure: Fig 8; 131pp; English.  
 PS  
 XX The sequence represents the amino acid sequence of synptosomal-  
 CC associated protein, SNAP25. The sequence was used to  
 CC create SNAP-25 double/single point mutants and C-terminal deletion  
 CC mutants used in a new method of treating a patient suffering from  
 CC poisoning or at risk of poisoning by a clostridial toxin, comprising  
 CC supplying a SNARE (soluble (N-ethylmaleimide-sensitive fusion protein) -  
 CC attachment protein receptor) to a cell of the patient, where the SNARE is  
 CC resistant to proteolysis by the toxin (toxin-resistant SNARE) and/or is  
 CC capable of inhibiting the toxin (toxin-inhibitory SNARE). The protein can  
 CC be used in a method of treating a patient in need of inhibition of SNARE-  
 CC dependent exocytosis from a cell capable of performing SNARE-dependent  
 CC exocytosis, comprising supplying a fragment, variant, fusion or derivative  
 CC of a SNARE or an inhibitory SNARE to the cell of the patient. The toxin  
 CC resistant or toxin inhibitory SNARE or a recombinant polynucleotide  
 CC encoding the SNARE is useful in the manufacture of a medicament for the  
 CC treatment of a patient suffering from poisoning or at risk of poisoning  
 CC by clostridial toxin, e.g. from botulism or tetanus. The fragment,  
 CC variant, fusion or derivative of a SNARE or of an inhibitory SNARE, or a  
 CC recombinant polynucleotide encoding either of these SNARE polypeptides  
 CC are useful in the manufacture of medicament for the treatment of a  
 CC patient in need of inhibition of SNARE-dependent exocytosis from a cell  
 CC capable of performing SNARE-dependent exocytosis. The method of treatment  
 CC is relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC  
 SQ Sequence 206 AA;  
 Query Match 100.0%; Score 1048; DB 22; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 6,7e-91;  
 Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVLEESKDAIGRTIVYALDEOGQQLERI 60  
 DB 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVLEESKDAIGRTIVYALDEOGQQLERI 60  
 QY 61 EEGMDQINKDKMEAEKKNITLDGKFCGLCYVCPCKNKLKSSDAYKKAGNNDGVAASOPARV 120  
 DB 61 EEGMDQINKDKMEAEKKNITLDGKFCGLCYVCPCKNKLKSSDAYKKAGNNDGVAASOPARV 120

QY 121 VDERQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
 |||||||  
 Db 121 VDERQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
 QY 181 IMEKADSNKTRIDEANQRATKMLGSG 206  
 |||||||  
 Db 181 IMEKADSNKTRIDEANQRATKMLGSG 206  
 RESULT 5  
 AAU00253  
 ID AAU00253 standard; Protein: 206 AA.  
 AC AAU00253;  
 XX  
 XX 12-SEP-2001 (first entry)  
 DE SNARE homologue, synaptosomal-associated protein, hSNAP25b.  
 XX  
 XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptosomal-associated protein; hSNAP25b; human.  
 XX  
 XX Homo sapiens.  
 OS  
 PN MO200118038-A2.  
 XX  
 XX 15-MAR-2001.  
 XX  
 XX 18-AUG-2000; 2000MO-GB03196.  
 XX  
 XX 20-AUG-1999; 99US-0149993.  
 XX  
 XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 PA Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
 PI  
 DR N-PSDB; AAS00370.  
 XX  
 PT Treating a patient suffering from poisoning or at risk of poisoning by  
 a clostridial toxin, e.g. botulism, comprises administering a  
 toxin-resistant or toxin-inhibitory SNARE -  
 XX  
 XX Disclosure; Fig 8; 130pp; English.  
 PS  
 XX The sequence represents the amino acid sequence of SNARE homologue,  
 CC synaptosomal-associated membrane protein, hSNAP25b, used during analysis  
 CC of SNAP-25. SNAP-25 mutants were used in a new method of treating a  
 CC patient suffering from poisoning or at risk of poisoning by a clostridial  
 CC toxin, comprising supplying a SNARE (soluble (N-ethylmaleimide-sensitive  
 CC fusion protein)-attachment protein receptor) to a cell of the patient,  
 CC where the SNARE is resistant to proteolysis by the toxin (toxin-resistant  
 CC SNARE) and/or is capable of inhibiting the toxin (toxin-inhibitory  
 CC SNARE). The protein can be used in a method of treating a patient in need  
 CC of inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis, comprising supplying a fragment,  
 CC variant, fusion or derivative of a SNARE or an inhibitory SNARE to the  
 CC cell of the patient. The toxin resistant or toxin inhibitory SNARE or a  
 CC recombinant polynucleotide encoding the SNARE is useful in the  
 CC manufacture of a medicament for the treatment of a patient suffering from  
 CC poisoning or at risk of poisoning by clostridial toxin, e.g. from  
 CC botulism or tetanus. The fragment, variant, fusion or derivative of a  
 CC SNARE or of an inhibitory SNARE, or a recombinant polynucleotide encoding  
 CC either of these SNARE polypeptides are useful in the manufacture of  
 CC medicament for the treatment of a patient in need of inhibition of SNARE-  
 CC dependent exocytosis from a cell capable of performing SNARE-dependent  
 CC exocytosis. The method of treatment is relatively fast, thus  
 CC alleviating the symptoms when most severe and taking the patient out of  
 CC critical state.  
 XX  
 XX Sequence 206 AA:  
 SQ

Query Match 100.0%; Score 1048; DB 22; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 6.7e-91;  
 Matches 206; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MAEDADMENELEEMQRRADQLADESLESTRRLQLVESKDAGITFLVLMDEQGLERI 60  
 |||||||  
 Db 1 MAEDADMENELEEMQRRADQLADESLESTRRLQLVESKDAGITFLVLMDEQGLERI 60  
 QY 61 EEGMDQINKDKMEAEKKNLTDLGKFCGLCYCPCKNLKSSDAIKKANGNODGVVASQPARV 120  
 |||||||  
 Db 61 EEGMDQINKDKMEAEKKNLTDLGKFCGLCYCPCKNLKSSDAIKKANGNODGVVASQPARV 120  
 QY 121 VDERQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
 |||||||  
 Db 121 VDERQMAISGGFIRRVYNDARENEMDENLEQVSGIIGNLRHMLDMGNEIDTQNRQIDR 180  
 QY 181 IMEKADSNKTRIDEANQRATKMLGSG 206  
 |||||||  
 Db 181 IMEKADSNKTRIDEANQRATKMLGSG 206  
 RESULT 6  
 AAU02640  
 ID AAU02640 standard; Protein: 206 AA.  
 AC AAU02640;  
 XX  
 XX 12-SEP-2001 (first entry)  
 DE Synaptosomal-associated protein, SNAP25, mutant L203A.  
 XX  
 XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptosomal-associated protein; mouse; mutant; mutein;  
 KM N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 XX  
 XX Mus sp.  
 OS  
 OS Synthetic.  
 OS  
 FH Key Location/Qualifiers  
 FT MISC-difference 203  
 FT /note= "Wild-type Leu substituted by Ala"  
 XX  
 XX MO200118038-A2.  
 XX  
 XX 15-MAR-2001.  
 XX  
 XX 18-AUG-2000; 2000MO-GB03196.  
 XX  
 XX 20-AUG-1999; 99US-0149993.  
 XX  
 XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 PA Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
 PI  
 DR N-PSDB; AAS00370.  
 XX  
 PT Treating a patient suffering from poisoning or at risk of poisoning by  
 a clostridial toxin, e.g. botulism, comprises administering a  
 toxin-resistant or toxin-inhibitory SNARE -  
 XX  
 XX Example 1; Page - : 131pp; English.  
 PS  
 XX The sequence represents the amino acid sequence of synaptosomal-  
 CC associated protein, SNAP25, mutant L203A, used in a new  
 CC method of treating a patient suffering from poisoning or at risk of  
 CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
 CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
 CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
 CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
 CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
 CC treating a patient in need of inhibition of SNARE-dependent exocytosis



CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
 CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
 CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
 CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
 CC useful in the manufacture of a medicament for the treatment of a patient  
 CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
 CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
 CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
 CC polynucleotide encoding either of these SNARE polypeptides are useful in  
 CC the manufacture of medicament for the treatment of a patient in need of  
 CC inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis. The method of treatment is  
 CC relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC Note: The present sequence is not shown in the specification but is  
 CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).  
 XX  
 SQ Sequence 206 AA;

Query Match 99.5%; Score 1043; DB 22; Length 206;  
 Best Local Similarity 99.5%; Pred. No. 2e-90; Indels 0; Gaps 0;  
 Matches 205; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVYESKDGAGIRTVMLDQEGQLERI 60  
 DB 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVYESKDGAGIRTVMLDQEGQLERI 60  
 QY 61 EEGMOQINKMDKEAEKNTLDGKFCGLCYPCPCNKLKSSDAYKKAGNNDGVVASQPARV 120  
 DB 61 EEGMOQINKMDKEAEKNTLDGKFCGLCYPCPCNKLKSSDAYKKAGNNDGVVASQPARV 120  
 QY 121 VDEROMATISGGFIRRVNDARENEMDENLEQVSGIIGLRLMALDMGNEIDTQNRQIDR 180  
 DB 121 VDEROMATISGGFIRRVNDARENEMDENLEQVSGIIGLRLMALDMGNEIDTQNRQIDR 180  
 QY 181 IMEKADSNKTRIDEANQATKMLGSG 206  
 DB 181 IMEKADSNKTRIDEANQATKMLGSG 206

RESULT 7  
 AAU00259  
 ID AAU00259 standard; Protein: 206 AA.  
 XX  
 AC AAU00259;  
 XX  
 DT 12-SEP-2001 (first entry)  
 DE Synaptoosomal-associated protein, SNAP25, mutant R198T.  
 XX  
 KM SNAP-25: poisoning; clostridial toxin; SNARE: toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptoosomal-associated protein; mouse; mutant; mutuin;  
 KM N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 OS Mus sp.  
 OS Synthetic.  
 XX  
 FT Key Location/Qualifiers  
 FT Misc-difference 198 /note="Wild-type Arg substituted by Thr"  
 XX  
 XX  
 PN MO200118038-A2.  
 XX  
 PD 15-MAR-2001.  
 XX  
 PF 18-AUG-2000; 2000MO-GB03196.  
 XX  
 PR 20-AUG-1999; 99US-0149993.  
 XX  
 PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 XX

PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
 XX  
 DR WPI, 2001-226739/23.  
 XX  
 PT Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -  
 XX  
 PS Example 1; Page - ; 131pp; English.  
 XX  
 XX The sequence represents the amino acid sequence of synaptoosomal-  
 CC associated protein, SNAP25, mutant R198T used in a new  
 CC method of treating a patient suffering from poisoning or at risk of  
 CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
 CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
 CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
 CC the toxin (toxin-inhibitory SNARE) and/or is capable of inhibiting the  
 CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
 CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
 CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
 CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
 CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
 CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
 CC useful in the manufacture of a medicament for the treatment of a patient  
 CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
 CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
 CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
 CC polynucleotide encoding either of these SNARE polypeptides are useful in  
 CC the manufacture of medicament for the treatment of a patient in need of  
 CC inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis. The method of treatment is  
 CC relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC Note: The present sequence is not shown in the specification but is  
 CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).  
 XX  
 SQ Sequence 206 AA;

Query Match 99.4%; Score 1042; DB 22; Length 206;  
 Best Local Similarity 99.5%; Pred. No. 2.5e-90;  
 Matches 205; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVYESKDGAGIRTVMLDQEGQLERI 60  
 DB 1 MAEDADMNELEEMORRADQDLADESLESTRMLQVYESKDGAGIRTVMLDQEGQLERI 60  
 QY 61 EEGMOQINKMDKEAEKNTLDGKFCGLCYPCPCNKLKSSDAYKKAGNNDGVVASQPARV 120  
 DB 61 EEGMOQINKMDKEAEKNTLDGKFCGLCYPCPCNKLKSSDAYKKAGNNDGVVASQPARV 120  
 QY 121 VDEROMATISGGFIRRVNDARENEMDENLEQVSGIIGLRLMALDMGNEIDTQNRQIDR 180  
 DB 121 VDEROMATISGGFIRRVNDARENEMDENLEQVSGIIGLRLMALDMGNEIDTQNRQIDR 180  
 QY 181 IMEKADSNKTRIDEANQATKMLGSG 206  
 DB 181 IMEKADSNKTRIDEANQATKMLGSG 206

RESULT 8  
 AAU00260  
 ID AAU00260 standard; Protein: 206 AA.  
 XX  
 AC AAU00260;  
 XX  
 DT 12-SEP-2001 (first entry)  
 DE Synaptoosomal-associated protein, SNAP25, mutant Q197A  
 XX  
 KM SNAP-25: poisoning; clostridial toxin; SNARE: toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptoosomal-associated protein; mouse; mutant; mutuin;  
 KM N-ethylmaleimide-sensitive fusion protein;

```

KM soluble NSF-attachment protein receptor.
XX
OS Mus sp.
OS Synthetic.
FH Key Location/Qualifiers
FT Misc-difference 197
FT /note="Wild-type Gln substituted by Ala"
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX
XX 20-AUG-1999; 99US-0149993.
XX
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
XX WPI; 2001-226739/23.
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - : 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant Q197A, used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis
XX from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant
XX polynucleotide encoding either of these SNARE polypeptides are useful in
XX the manufacture of medicament for the treatment of a patient in need of
XX inhibition of SNARE-dependent exocytosis. The method of treatment is
XX relatively fast, thus alleviating the symptoms when most severe and
XX taking the patient out of critical state.
XX Note: The present sequence is not shown in the specification but is
XX derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX
XX Sequence 206 AA:
XX
XX Query Match 99.4%; Score 1042; DB 22; Length 206;
XX Best Local Similarity 99.5%; Pred. No. 2,5e-90;
XX Matches 205; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

OY 181 IMEKADSNKTRIDEANORATKMLGSG 206
XX
XX
XX Db 181 IMEKADSNKTRIDEANARATKMLGSG 206
XX
XX
XX RESULT 9
XX AAU00261
XX ID AAU00261 standard; Protein; 206 AA.
XX
XX AAU00261;
XX
XX 12-SEP-2001 (first entry)
XX
XX Synaptosomal-associated protein, SNAP25, mutant R198A.
XX
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
XX synaptosomal-associated protein; mouse; mutant; mutein;
XX N-ethylmaleimide-sensitive fusion protein;
XX soluble NSF-attachment protein receptor.
XX
XX Mus sp.
XX Synthetic.
XX
XX Key Location/Qualifiers
XX FT Misc-difference 198
XX FT /note="Wild-type Arg substituted by Ala"
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX
XX 20-AUG-1999; 99US-0149993.
XX
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
XX WPI; 2001-226739/23.
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - : 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant R198A used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis
XX from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant
XX polynucleotide encoding either of these SNARE polypeptides are useful in
XX the manufacture of medicament for the treatment of a patient in need of
XX inhibition of SNARE-dependent exocytosis. The method of treatment is
XX relatively fast, thus alleviating the symptoms when most severe and
XX taking the patient out of critical state.
XX Note: The present sequence is not shown in the specification but is
XX derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX

```

```

XX Sequence      206 AA:
SQ
Query Match      99.4%: Score 1042; DB 22; Length 206;
Best Local Similarity 99.5%: Pred. No. 2.5e-90;
Matches 205; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 1 MAEDADMNLEEMORRRADQLADESLESTRRLQLVEESKDAGIRTLVW.DBOGQLERI 60
DB 1 MAEDADMNLEEMORRRADQLADESLESTRRLQLVEESKDAGIRTLVW.DBOGQLERI 60
OY 61 EEGMDQINKDKMEAEKNLTDLGKFCGLCYPCPNKLSSDAYKKANGNNDGVVASQPARV 120
DB 61 EEGMDQINKDKMEAEKNLTDLGKFCGLCYPCPNKLSSDAYKKANGNNDGVVASQPARV 120
OY 121 VDERQMAISGCFIRRYVNDARENEMDENLEQVSGIIGLRHMLDMGNEIDTONRQIDR 180
DB 121 VDERQMAISGCFIRRYVNDARENEMDENLEQVSGIIGLRHMLDMGNEIDTONRQIDR 180
OY 181 IMEKADSNKTRIDEANORATKMLGSG 206
DB 181 IMEKADSNKTRIDEANORATKMLGSG 206

RESULT 10
AAU02638
ID AAU02638 standard; Protein: 206 AA.
XX
AC AAU02638:
XX
DT 12-SEP-2001 (first entry)
DE Synaptosomal-associated protein, SNAP25, mutant M202A.
XX
KW SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
KW synaptosomal-associated protein; mouse; mutant; mutelin;
KW N-ethylmaleimide-sensitive fusion protein;
XX soluble NSF-attachment protein receptor.
XX
OS Mus sp.
OS Synthetic.
XX
FH Key Location/Qualifiers
FT Misc-difference 202 /note= "Wild-type Met substituted by Ala"
FT
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX
XX 20-AUG-1999; 99US-0149993.
XX
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
XX DOLLY JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
XX WPI: 2001-226739/23.
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - ; 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant M202A, used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX

```

```

CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of
CC treating a patient in need of inhibition of SNARE-dependent exocytosis
CC from a cell capable of performing SNARE-dependent exocytosis, comprises
CC supplying a fragment, variant, fusion or derivative of a SNARE or an
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
CC useful in the manufacture of a medicament for the treatment of a patient
CC suffering from poisoning or at risk of poisoning by clostridial toxin,
CC e.g. from botulism or tetanus. The fragment, variant, fusion or
CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant
CC polynucleotide encoding either of these SNARE polypeptides are useful in
CC the manufacture of medicament for the treatment of a patient in need of
CC inhibition of SNARE-dependent exocytosis from a cell capable of
CC performing SNARE-dependent exocytosis. The method of treatment is
CC relatively fast, thus alleviating the symptoms when most severe and
CC taking the patient out of critical state.
CC Note: The present sequence is not shown in the specification but is
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU0246).
XX
SQ Sequence      206 AA:
Query Match      99.4%: Score 1042; DB 22; Length 206;
Best Local Similarity 99.5%: Pred. No. 2.5e-90;
Matches 205; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 1 MAEDADMNLEEMORRRADQLADESLESTRRLQLVEESKDAGIRTLVW.DBOGQLERI 60
DB 1 MAEDADMNLEEMORRRADQLADESLESTRRLQLVEESKDAGIRTLVW.DBOGQLERI 60
OY 61 EEGMDQINKDKMEAEKNLTDLGKFCGLCYPCPNKLSSDAYKKANGNNDGVVASQPARV 120
DB 61 EEGMDQINKDKMEAEKNLTDLGKFCGLCYPCPNKLSSDAYKKANGNNDGVVASQPARV 120
OY 121 VDERQMAISGCFIRRYVNDARENEMDENLEQVSGIIGLRHMLDMGNEIDTONRQIDR 180
DB 121 VDERQMAISGCFIRRYVNDARENEMDENLEQVSGIIGLRHMLDMGNEIDTONRQIDR 180
OY 181 IMEKADSNKTRIDEANORATKMLGSG 206
DB 181 IMEKADSNKTRIDEANORATKMLGSG 206

RESULT 11
AAU0262
ID AAU0262 standard; Protein: 206 AA.
XX
AC AAU0262:
XX
DT 12-SEP-2001 (first entry)
DE Synaptosomal-associated protein, SNAP25, mutant Q197K, R198K.
XX
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
XX synaptosomal-associated protein; mouse; mutant; mutelin;
XX N-ethylmaleimide-sensitive fusion protein;
XX soluble NSF-attachment protein receptor.
XX
XX Mus sp.
XX Synthetic.
XX
FH Key Location/Qualifiers
FT Misc-difference 197 /note= "Wild-type Gln substituted by 'lys'"
FT Misc-difference 198 /note= "Wild-type Arg substituted by 'lys'"
FT
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX

```

```

XX 20-AUG-1999; 99US-0149993.
PR (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
PA Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX MPI: 2001-226739/23.
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - : 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant Q197K/R198K, used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis
XX from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant
XX polynucleotide encoding either of these SNARE polypeptides are useful in
XX the manufacture of medicament for the treatment of a patient in need of
XX inhibition of SNARE-dependent exocytosis from a cell capable of
XX performing SNARE-dependent exocytosis. The method of treatment is
XX relatively fast, thus alleviating the symptoms when most severe and
XX taking the patient out of critical state.
XX Note: The present sequence is not shown in the specification but is
XX derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX
XX Sequence 206 AA:
XX
XX Query Match 99.3%; Score 1041; DB 22; Length 206;
XX Best Local Similarity 99.0%; Pred. No. 3.1e-90;
XX Matches 204; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
XX
XX QY 1 MAEDADMRNELEEMQRRADQLADESLESTRMQLVYESKQAGIRITVMDDEGEQLERI 60
XX DB 1 MAEDADMRNELEEMQRRADQLADESLESTRMQLVYESKQAGIRITVMDDEGEQLERI 60
XX QY 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKKMGNNQDGVASQPARV 120
XX DB 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKKMGNNQDGVASQPARV 120
XX QY 121 VDEREQMAISGGFIRRYVNDARENEMDENLEQVSGIIGNLRHMAIDMGNEIDTONRQIDR 180
XX DB 121 VDEREQMAISGGFIRRYVNDARENEMDENLEQVSGIIGNLRHMAIDMGNEIDTONRQIDR 180
XX QY 181 IMEKADSNKTRIDEANQATKMLGSG 206
XX DB 181 IMEKADSNKTRIDEANQATKMLGSG 206
XX
XX RESULT 12
XX AAU00257
XX ID AAU00257 standard; Protein: 206 AA.
XX
XX AC AAU00257;
XX
XX 12-SEP-2001 (first entry)
XX
XX Synaptosomal-associated protein, SNAP25, mutant Q197A/R198K.

```

```

XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
XX synaptosomal-associated protein; mouse; mutant; mutein;
XX N-ethylmaleimide-sensitive fusion protein;
XX soluble NSF-attachment protein receptor.
XX
XX Mus sp.
XX Synthetic.
XX
XX Key Location/Qualifiers
XX Misc-difference 197
XX Misc-difference 198
XX Misc-difference 198
XX Note="Wild-type Arg substituted by Lys"
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX
XX 20-AUG-1999; 99US-0149993.
XX
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX MPI: 2001-226739/23.
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - : 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant Q197A/R198K, used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis
XX from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant
XX polynucleotide encoding either of these SNARE polypeptides are useful in
XX the manufacture of medicament for the treatment of a patient in need of
XX inhibition of SNARE-dependent exocytosis from a cell capable of
XX performing SNARE-dependent exocytosis. The method of treatment is
XX relatively fast, thus alleviating the symptoms when most severe and
XX taking the patient out of critical state.
XX Note: The present sequence is not shown in the specification but is
XX derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX
XX Sequence 206 AA:
XX
XX Query Match 99.1%; Score 1039; DB 22; Length 206;
XX Best Local Similarity 99.0%; Pred. No. 4.8e-90;
XX Matches 204; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
XX
XX QY 1 MAEDADMRNELEEMQRRADQLADESLESTRMQLVYESKQAGIRITVMDDEGEQLERI 60
XX DB 1 MAEDADMRNELEEMQRRADQLADESLESTRMQLVYESKQAGIRITVMDDEGEQLERI 60
XX QY 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKKMGNNQDGVASQPARV 120

```

```

|||||
Db 61 EEGMDQINKDKAEKKNLTDLGKFCGLCCPCNKLKSSDAYKKAGNNODGVVASQPARV 120
Qy 121 VDEREQMAISGCFIRRYTNDARENEMDENLEQVSGIIGLRHMAIDMNEIDTQNRQIDR 180
Db 121 VDEREQMAISGCFIRRYTNDARENEMDENLEQVSGIIGLRHMAIDMNEIDTQNRQIDR 180
Qy 181 IMEKADSNKTRIDEANQRTKMLGSG 206
Db 181 IMEKADSNKTRIDEANQRTKMLGSG 206

RESULT 13
AAU00266
ID AAU00266 standard; Protein: 206 AA.
XX
AC AAU00266;
XX
DT 12-SEP-2001 (first entry)
XX
DE Synaptosomal-associated protein, SNAP25, mutant Q197K/R198H.
KM SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
KM synaptosomal-associated protein; mouse; mutant; mutant;
KM N-ethylmaleimide-sensitive fusion protein;
KM soluble NSF-attachment protein receptor.
XX
OS Mus sp.
OS Synthetic.
XX
FH Key Location/Qualifiers
FT Misc-difference 197 /note= "Wild-type Gln substituted by Lys"
FT Misc-difference 198 /note= "Wild-type Arg substituted by His"
FT MO200118038-A2.
XX
PD 15-MAR-2001.
XX
PF 18-AUG-2000; 2000WO-GB03196.
XX
PR 20-AUG-1999; 99US-0149993.
XX
PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
DR WPI: 2001-226739/23.
XX
PT Treating a patient suffering from poisoning or at risk of poisoning by
PT a clostridial toxin, e.g. botulism, comprises administering a
PT toxin-resistant or toxin-inhibitory SNARE -
XX
PS Example 1: Page - : 131pp; English.
XX
CC The sequence represents the amino acid sequence of synaptosomal-
CC associated protein, SNAP25, mutant Q197K/R198H, used in a new
CC method of treating a patient suffering from poisoning or at risk of
CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
CC to a cell of the patient, where the SNARE is resistant to proteolysis by
CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of
CC treating a patient in need of inhibition of SNARE-dependent exocytosis
CC from a cell capable of performing SNARE-dependent exocytosis, compris-
CC supplying a fragment, variant, fusion or derivative of a SNARE or an
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
CC useful in the manufacture of a medicament for the treatment of a patient
CC suffering from poisoning or at risk of poisoning by clostridial toxin,
CC e.g. from botulism or tetanus. The fragment, variant, fusion or

```

```

CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant
CC polynucleotide encoding either of these SNARE polypeptides are useful in
CC the manufacture of medicament for the treatment of a patient in need of
CC inhibition of SNARE-dependent exocytosis from a cell capable of
CC performing SNARE-dependent exocytosis. The method of treatment is
CC relatively fast, thus alleviating the symptoms when most severe and
CC taking the patient out of critical state.
CC Note: The present sequence is not shown in the specification but is
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX
SQ Sequence 206 AA;
XX
Query Match 99.1%; Score 1039; DB 22; Length 206;
Best Local Similarity 99.0%; Pred. No. 4.8e-90;
Matches 204; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
Qy 1 MAEDADMNLEEMQRRADQADESLESTRMLQLVEESKAGITLVYALDEQEQLERI 60
Db 1 MAEDADMNLEEMQRRADQADESLESTRMLQLVEESKAGITLVYALDEQEQLERI 60
Qy 61 EEGMDQINKDKAEKKNLTDLGKFCGLCCPCNKLKSSDAYKKAGNNODGVVASQPARV 120
Db 61 EEGMDQINKDKAEKKNLTDLGKFCGLCCPCNKLKSSDAYKKAGNNODGVVASQPARV 120
Qy 121 VDEREQMAISGCFIRRYTNDARENEMDENLEQVSGIIGLRHMAIDMNEIDTQNRQIDR 180
Db 121 VDEREQMAISGCFIRRYTNDARENEMDENLEQVSGIIGLRHMAIDMNEIDTQNRQIDR 180
Qy 181 IMEKADSNKTRIDEANQRTKMLGSG 206
Db 181 IMEKADSNKTRIDEANQRTKMLGSG 206

RESULT 14
AAU02171
ID AAU02171 standard; Protein: 206 AA.
XX
AC AAU02171;
XX
DT 12-SEP-2001 (first entry)
XX
DE Synaptosomal-associated protein, SNAP25, mutant R198Y/L203A.
XX
KM SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
KM synaptosomal-associated protein; mouse; mutant; mutant;
KM N-ethylmaleimide-sensitive fusion protein;
KM soluble NSF-attachment protein receptor.
XX
OS Mus sp.
OS Synthetic.
XX
FH Key Location/Qualifiers
FT Misc-difference 198 /note= "Wild-type Arg substituted by Thr"
FT Misc-difference 203 /note= "Wild-type Leu substituted by Ala"
FT MO200118038-A2.
XX
PD 15-MAR-2001.
XX
PF 18-AUG-2000; 2000WO-GB03196.
XX
PR 20-AUG-1999; 99US-0149993.
XX
PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
DR WPI: 2001-226739/23.
XX
PT Treating a patient suffering from poisoning or at risk of poisoning by

```

PT a clostridial toxin, e.g. botulism, comprises administering a  
PT toxin-resistant or toxin-inhibitory SNARE -  
PS Example 1; Page - ; 131pp; English.

XX The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25, mutant R198T/L203A, used in a new  
CC method of treating a patient suffering from poisoning or at risk of  
CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
CC useful in the manufacture of a medicament for the treatment of a patient  
CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
CC polynucleotide encoding either of these SNARE polypeptides are useful in  
CC the manufacture of medicament for the treatment of a patient in need of  
CC inhibition of SNARE-dependent exocytosis. The method of treatment is  
CC relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
CC Note: The present sequence is not shown in the specification but is  
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).

XX Sequence 206 AA:

Query Match 99.0%; Score 1037; DB 22; Length 206;  
Best Local Similarity 99.0%; Pred. No. 7.3e-90;  
Matches 204; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 MAEDADNRNELEEMQRRADOLADESLESTRMQLVVEESKAGIRITVLMDEOGEOLERI 60  
DB 1 MAEDADNRNELEEMQRRADOLADESLESTRMQLVVEESKAGIRITVLMDEOGEOLERI 60  
QY 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKAMGNODGVVASOPARV 120  
DB 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKAMGNODGVVASOPARV 120  
QY 121 VDREQMAISGCFIRRYTNDARENMENDELEQVSGIIGNLRHMLDGMNEIDTONRQIDR 180  
DB 121 VDREQMAISGCFIRRYTNDARENMENDELEQVSGIIGNLRHMLDGMNEIDTONRQIDR 180  
QY 181 IMERKADSNKTRIDEANQATKMLGSG 206  
DB 181 IMERKADSNKTRIDEANQATKMLGSG 206

RESULT 15  
AAU00256  
ID AAU00256 standard; Protein: 206 AA.

XX AAU00256:  
DT 12-SEP-2001 (first entry)

XX Synaptosomal-associated protein, SNAP25, mutant Q197A/R198A.

XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
KM synaptosomal-associated protein; mouse; mutant; mutein;  
KM N-ethylmaleimide-sensitive fusion protein;  
KM soluble NSF-attachment protein receptor.

XX Mus sp.  
OS Synthetic.

XX

FH Key Location/Qualifiers  
FT Misc-difference 197  
FT /note= "Wild-type Gln substituted by Ala"  
FT Misc-difference 198  
FT /note= "Wild-type Arg substituted by Ala"

XX WO200118038-A2.

XX 15-MAR-2001.

XX 18-AUG-2000; 2000WO-GB03196.

XX 20-AUG-1999; 99US-0149993.

XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.

XX DOLLY JO, O'Sullivan GA, Mohammed N, Foran PG;

XX WPI; 2001-226739/23.

XX Treating a patient suffering from poisoning or at risk of poisoning by  
PT a clostridial toxin, e.g. botulism, comprises administering a  
PT toxin-resistant or toxin-inhibitory SNARE -

XX Example 1; Page - ; 131pp; English.

XX The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25, mutant Q197A/R198A, used in a new  
CC method of treating a patient suffering from poisoning or at risk of  
CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
CC useful in the manufacture of a medicament for the treatment of a patient  
CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
CC polynucleotide encoding either of these SNARE polypeptides are useful in  
CC the manufacture of medicament for the treatment of a patient in need of  
CC inhibition of SNARE-dependent exocytosis. The method of treatment is  
CC relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
CC Note: The present sequence is not shown in the specification but is  
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).

XX Sequence 206 AA:

Query Match 98.9%; Score 1036; DB 22; Length 206;  
Best Local Similarity 99.0%; Pred. No. 9.1e-90;  
Matches 204; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 MAEDADNRNELEEMQRRADOLADESLESTRMQLVVEESKAGIRITVLMDEOGEOLERI 60

DB 1 MAEDADNRNELEEMQRRADOLADESLESTRMQLVVEESKAGIRITVLMDEOGEOLERI 60

QY 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKAMGNODGVVASOPARV 120

DB 61 EEGMDQINKDKKEAEKNTLDGKFCGCLVCPCKNLKSSDAKKAMGNODGVVASOPARV 120

QY 121 VDREQMAISGCFIRRYTNDARENMENDELEQVSGIIGNLRHMLDGMNEIDTONRQIDR 180

DB 121 VDREQMAISGCFIRRYTNDARENMENDELEQVSGIIGNLRHMLDGMNEIDTONRQIDR 180

QY 181 IMERKADSNKTRIDEANQATKMLGSG 206

DB 181 IMERKADSNKTRIDEANQATKMLGSG 206

Tue Dec 3 08:36:05 2002

pct-us02-27145-2.rag

Page 11

Search completed: November 19, 2002, 17:34:26  
Job time : 87.2973 secs

---

**THIS PAGE BLANK (USP10)**



GenCore version 5.1.3  
Copyright (c) 1993 - 2002 Compugen Ltd.

## OM protein - protein search, using sw model

Run on: November 19, 2002, 17:32:44 : Search time 3.3515 Seconds  
(without alignments)  
318.082 Million cell updates/sec

Title: PCT-US02-27145-8  
Perfect score: 39  
Sequence: 1 QIDRIMEK 8

Scoring table: BIOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues  
Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database :  
1: A\_Geneseq\_101002: \*  
2: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1980.DAT: \*  
3: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1981.DAT: \*  
4: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1982.DAT: \*  
5: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1983.DAT: \*  
6: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1984.DAT: \*  
7: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1985.DAT: \*  
8: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1986.DAT: \*  
9: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1987.DAT: \*  
10: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1988.DAT: \*  
11: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1989.DAT: \*  
12: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1990.DAT: \*  
13: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1991.DAT: \*  
14: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1992.DAT: \*  
15: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1993.DAT: \*  
16: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1994.DAT: \*  
17: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1995.DAT: \*  
18: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1996.DAT: \*  
19: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1997.DAT: \*  
20: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1998.DAT: \*  
21: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA1999.DAT: \*  
22: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2000.DAT: \*  
23: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2001.DAT: \*  
24: /SIDS2/gcgdata/geneseq/geneseq-emb1/AA2002.DAT: \*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	39	100.0	20	AAW30098	Neurotransmitter s
2	39	100.0	37	AAW30097	Neurotransmitter s
3	39	100.0	49	AAW5386	Human brain expres
4	39	100.0	70	AAW8623	SNAP-25 residues 1
5	39	100.0	86	AAW15584	Human SNAP-25 N-te
6	39	100.0	116	AAW15165	Clostridial neurot
7	39	100.0	116	AAW15166	Clostridial neurot
8	39	100.0	198	AAW00255	Synaptosomal-assoc
9	39	100.0	199	AAW00263	Synaptosomal-assoc
10	39	100.0	200	AAW00264	Synaptosomal-assoc

11	39	100.0	201	AAW02637	Synaptosomal-assoc
12	39	100.0	202	AAW00265	Synaptosomal-assoc
13	39	100.0	203	AAW02636	Synaptosomal-assoc
14	39	100.0	206	AAW30103	Synaptosomal-assoc
15	39	100.0	206	AAW79198	House SNAP-25 poly
16	39	100.0	206	AAW43426	House synaptosomal
17	39	100.0	206	AAW00246	Synaptosomal-assoc
18	39	100.0	206	AAW00252	SNARE homologue, s
19	39	100.0	206	AAW00253	SNARE homologue, s
20	39	100.0	206	AAW00256	Synaptosomal-assoc
21	39	100.0	206	AAW00257	Synaptosomal-assoc
22	39	100.0	206	AAW00258	Synaptosomal-assoc
23	39	100.0	206	AAW00259	Synaptosomal-assoc
24	39	100.0	206	AAW00260	Synaptosomal-assoc
25	39	100.0	206	AAW00261	Synaptosomal-assoc
26	39	100.0	206	AAW00262	Synaptosomal-assoc
27	39	100.0	206	AAW00266	Synaptosomal-assoc
28	39	100.0	206	AAW002171	Synaptosomal-assoc
29	39	100.0	206	AAW02638	Synaptosomal-assoc
30	39	100.0	206	AAW02639	Synaptosomal-assoc
31	39	100.0	206	AAW02640	Synaptosomal-assoc
32	34	87.2	200	AAW82046	S. epidermidis ope
33	34	87.2	200	AAW82048	S. epidermidis ope
34	34	87.2	208	ABP38940	Staphylococcus epi
35	33	84.6	84.1	ABG09845	Novel human diagno
36	33	84.6	164.3	ABG63371	Drosophila melano
37	33	84.6	182.5	ABG09849	Novel human diagno
38	33	84.6	1909	ABG21157	Novel human diagno
39	32	82.1	35	AAW78070	Influenza strain A
40	32	82.1	566	AAW29746	Influenza virus ha
41	31	79.5	59	AAW02582	Human polypeptide
42	31	79.5	135	AAW36760	Amino acid sequenc
43	31	79.5	349	ABG01597	Novel human diagno
44	31	79.5	429	AAW44050	Arabidopsis thalia
45	31	79.5	429	AAW51986	Arabidopsis thalia

## ALIGNMENTS

RESULT 1	
AAW30098	AAW30098 standard; peptide: 20 AA.
ID	
AC	AAW30098;
XX	
DT	06-APR-1998 (first entry)
XX	
DE	Neurotransmitter secretion inhibitor #2.
XX	
KW	Neurotransmitter secretion; neuronal cell; synaptic vesicle;
KW	excitation-secretory uncoupling peptide; catecholamine secretion;
KW	bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;
KW	synaptosomal associated protein; SNAP-25.
XX	
OS	Homo sapiens.
XX	
PN	W09734620-A1.
XX	
PD	25-SEP-1997.
XX	
PF	18-MAR-1997; 97W0-US04393.
XX	
PR	18-MAR-1996; 96US-0013599.
XX	
PA	(REGC) UNIV CALIFORNIA.
XX	
PI	Montal M;
XX	
DR	WPI: 1997-479986/44.
XX	
PT	Excitation-secretory uncoupling peptide(s) for inhibiting
PT	neurotransmitter release - used particularly for treating muscle

```

PT Spasticity, and for delivering drugs specifically to neural cells
XX
PS Claim 12; Page 31; 61pp; English.
XX
CC This sequence corresponds to residues 170-189 of the human 25 kD
CC synaptosomal associated protein (SNAP-25), and is a inhibitory agent of
CC the invention. The agents of the invention inhibit secretion of
CC neurotransmitter from neuronal cells and is an excitation-secretory
CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which
CC correspond substantially to any one of AAW30097-W30102, or more
CC generally any (I) that inhibits 50% of catecholamine secretion from
CC bovine chromaffin cells at a concentration of 10 microm, especially 0.25
CC microm, or less. (I) are used, as a replacement for Clostridium toxin, to
CC inhibit release of neurotransmitters from synaptic vesicles, specifically
CC for reducing muscle spasticity. Also (I) may be labelled to allow in
CC vivo imaging of intracellular distribution of (I). Compounds for
CC delivering the drug to neural cells provide targeted drug delivery, e.g.
CC of substance P to brain tumours for induction of apoptosis. Unlike the
CC neurotoxins, (I) are not toxic or immunogenic and are more readily
CC available. Their therapeutic effect lasts for several days or weeks, so
CC lower doses or less frequent treatments are required.
XX
XX Sequence 20 AA:
S0
XX
XX Query Match 100.0%; Score 39; DB 18; Length 20;
XX Best Local Similarity 100.0%; Pred. No. 0.4;
XX Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0.
OY 1 QIDRIMER 8
DB 8 QIDRIMER 15
XX
XX RESULT 2
XX AAW30097
XX ID AAW30097 standard; peptide: 37 AA.
XX
XX AAW30097:
XX
XX 06-APR-1998 (first entry)
XX
XX Neurotransmitter secretion inhibitor #1.
XX
XX Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;
XX excitation-secretory uncoupling peptide; catecholamine secretion;
XX bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;
XX synaptosomal associated protein; SNAP-25.
XX
XX Homo sapiens.
XX
XX WO9734620-A1.
XX
XX 25-SEP-1997.
XX
XX 18-MAR-1997; 97WO-US04393.
XX
XX 18-MAR-1996; 96US-0013599.
XX
XX (REGC ) UNIV CALIFORNIA.
XX
XX Montal M;
XX
XX MPI; 1997-479986/44.
XX
XX
XX Excitation-secretory uncoupling peptide(s) for inhibiting
XX neuro-transmitter release - used particularly for treating muscle
XX spasticity, and for delivering drugs specifically to neural cells
XX
XX Claim 1; Page 30; 61pp; English.
XX
XX This sequence corresponds to residues 170-206 of the human 25 kD
XX synaptosomal associated protein (SNAP-25), and is a inhibitory agent of
XX the invention. The agents of the invention inhibit secretion of

```

CC	neurotransmitter from neuronal cells and is an excitation-secretory
CC	uncoupling peptide (I) of at least 20 amino acids (aa) all of which
CC	correspond substantially to any one of AAM30097-W30102, or more
CC	generally any (I) that inhibits 50% of catecholamine secretion from
CC	bovine chromaffin cells at a concentration of 10 microm, especially 0.25
CC	microm, or less. (I) are used, as a replacement for Clostridium toxin, to
CC	inhibit release of neurotransmitters from synaptic vesicles, specifically
CC	for reducing muscle spasticity. Also (I) may be labelled to allow in
CC	vivo imaging of intracellular distribution of (I). Compounds for
CC	delivering the drug to neural cells provide targeted drug delivery, e.g.
CC	of substance P to brain tumours for induction of apoptosis. Unlike the
CC	neurotoxins, (I) are not toxic or immunogenic and are more readily
CC	available. Their therapeutic effect lasts for several days or weeks, so
CC	lower doses or less frequent treatments are required.
XX	
XX	Sequence 37 AA:
XX	
XX	Query Match 100.0%; Score 39; DB 18; Length 37;
XX	Best Local Similarity 100.0%; Pred. No. 0.73;
XX	Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
OY	1 QIDRIMER 8
DB	8 QIDRIMER 15
XX	
XX	AAAM57386 standard; Protein: 49 AA.
XX	
XX	AAAM57386;
XX	
XX	05-NOV-2001 (first entry)
XX	
XX	Human brain expressed single exon probe encoded protein SEQ ID NO: 29491.
XX	
XX	Human: brain expressed exon: gene expression analysis; probe:
XX	microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
XX	epilepsy; cancer.
XX	
XX	Homo sapiens.
XX	
XX	WO200157275-A2.
XX	
XX	09-AUG-2001.
XX	
XX	30-JAN-2001; 2001WO-US00667.
XX	
XX	04-FEB-2000; 2000US-0180312.
XX	26-MAY-2000; 2000US-0207456.
XX	30-JUN-2000; 2000US-0608408.
XX	03-AUG-2000; 2000US-0632366.
XX	21-SEP-2000; 2000US-0234687.
XX	27-SEP-2000; 2000US-0236359.
XX	04-OCT-2000; 2000GB-0024263.
XX	
XX	(MOLE-) MOLECULAR DYNAMICS INC.
XX	
XX	Penn SG, Hanzel DK, Chen W, Rank DR;
XX	
XX	WPI; 2001-483446/52.
XX	
XX	Single exon nucleic acid probes for analyzing gene expression in human
XX	brains -
XX	
XX	Example 4; SEQ ID NO: 29491; 650bp + Sequence Listing; English.
XX	
XX	The present invention provides a number of single exon nucleic acid
XX	probes which are derived from genomic sequences expressed in the human
XX	brain. They can be used to measure gene expression in brain cell samples,
XX	which may enable the diagnosis and improved treatment of nervous system
XX	diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia,
XX	epilepsy and cancers. The present sequence is a protein encoded by one of

CC the probes of the invention.

XX Sequence 49 AA;

Query Match 100.0%; Score 39; DB 22; Length 49;  
Best Local Similarity 100.0%; Pred. No. 0.96;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QIDRIMEX 8  
| | | | | | | |  
DB 41 QIDRIMEX 48

RESULT 4  
AAR86823  
ID AAR86823 standard; Peptide; 70 AA.

XX AAR86823;

XX 15-AUG-1996 (first entry)

XX SNAP-25 residues 137-206.

XX VAMP: vesicle-associated membrane protein; SNAP-25; syntaxin;  
KW neurotransmitter; neurotoxin; botulinum; botulism; cleavage;  
XX substrate; antibody; detection; assay.

XX Synthetic.

XX WO9533850-A1.

XX 14-DEC-1995.

XX 02-JUN-1995; 95MO-GB01279.

XX 03-JUN-1994; 94GB-0011138.

XX (CAMR-) CAMR CENT APPLIED MICROBIOLOGY & RES.

XX (MICR-) MICROBIOLOGICAL RES AUTHORITY.

XX Hallis B, James BAF, Quinn CP, Shone CC;

XX WPI; 1996-040249/04.

XX Assay for botulinum or tetanus toxin - by combining test cpd. with  
PT substrate which is cleaved by the toxin, and antibody specific for  
PT the cleaved but not uncleaved substrate

XX Example 4; Page 19; 48pp; English.

XX The botulinum neurotoxins possess highly specific zinc-endopeptidase  
CC activities within their light sub-units. Depending on the neurotoxin  
CC type these act to cleave small proteins within the nerve cell which are  
CC involved in neurotransmitter release. Antibodies are used in assays  
CC which detect cleaved but not uncleaved substrate. Assays for botulinum  
CC types A and E use the present sequence as a substrate. The sequence is  
CC SNAP-25 protein, residues 137-206.

XX Sequence 70 AA;

Query Match 100.0%; Score 39; DB 17; Length 70;  
Best Local Similarity 100.0%; Pred. No. 1.4;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QIDRIMEX 8  
| | | | | | | |  
DB 41 QIDRIMEX 48

RESULT 5  
AAB15584  
ID AAB15584 standard; peptide; 86 AA.

AC AAB15584;

XX 02-MAR-2001 (first entry)

XX Human SNAP-25 N-terminal peptide #4.

XX Dermatological; neuroprotective; relaxant; neuronal exocytosis modulator;  
KW SNAP-25; synaptosomal-associated protein 25; facial wrinkle; asymmetry;  
XX neurodegenerative disorder.

XX Homo sapiens.

XX WO200064932-A1.

XX 02-NOV-2000.

XX 18-FEB-2000; 2000MO-ES00058.

XX 23-APR-1999; 99ES-0000844.

XX (LIPOTEC) LIPOTEC SA.

XX Blanes Mira MC, Llobregat Hernandez MM, Gil Tebar AI;

XX Fernandez Bailester GU, Planell Cases RM, Ferrer Montiel AV;

XX Viniegra Bover S, Gutierrez Perez LM, Carbonell Castej T;

XX WPI; 2001-007091/01.

XX New peptides containing amino acid sequences from known proteins for  
PT treatment of neurological disorders

XX Claim 16; Page 32-33; 40pp; Spanish.

XX The invention relates to new peptides comprising 3-30 contiguous amino  
CC acids from the N-terminus of the protein SNAP-25

CC (synaptosomal-associated protein 25). The peptides AAB15581-B15586

CC represent examples of the peptides of the invention. The peptides have

CC neuronal exocytosis inhibitory activity and are used for: treatment of

CC facial wrinkles and asymmetry and pathological neuronal

CC exocytosis-mediated pathological disorders and alterations manifested

CC e.g. by spasms and neurological and neurodegenerative disorders.

XX Sequence 86 AA;

QY 1 QIDRIMEX 8  
| | | | | | | |  
DB 57 QIDRIMEX 64

RESULT 6  
AAB15165  
ID AAB15165 standard; peptide; 116 AA.

XX AAB15165;

XX 02-SEP-2002 (first entry)

XX Clostridial neurotoxin protease substrate peptide 4.

XX Clostridial neurotoxin substrate; botulinum neurotoxin substrate; FRET;

XX fluorescence resonant energy transfer assay; quenched-signal;

XX clostridial neurotoxin detection; food.

XX Unidentified.

XX key Location/Qualifiers

FT Modified-site 1 /note="S-fluoresceinyl-cysteine"

FT Cleavage-site 89..90  
FT /note= "The peptide is cleaved between these two  
FT residues by type E Clostridium botulinum neurotoxin"  
FT Cleavage-site 106..107  
FT /note= "The peptide is cleaved between these two  
FT residues by type A Clostridium botulinum neurotoxin"  
XX  
PN WO200225284-A2.  
XX  
PD 28-MAR-2002.  
XX  
PE 25-SEP-2001; 2001WO-US30188.  
XX  
PR 25-SEP-2000; 2000US-235050P.  
XX  
PA (USME-) US MEDICAL RES INST INFECTIOUS DISEASES.  
PI Schmidt JI, Stafford RG;  
XX  
XX WPI: 2002-499829/53.  
XX  
XX Substrate useful in e.g. an assay for the protease activity of  
PT clostridial neurotoxin, comprises modified peptide or protein -  
PT  
XX  
XX Claim 22; Page 16; 48pp; English.  
XX  
XX The invention comprises clostridial neurotoxin substrate peptides which  
CC can serve as fluorescence resonant energy transfer assay (FRET) or  
CC quenched-signal substrates in assays for the proteolytic activities of  
CC clostridial neurotoxins. The invention further comprises Clostridium  
CC botulinum neurotoxin substrate peptides that can serve as immobilised  
CC substrates (i.e. bound to a solid phase) in assays for the proteolytic  
CC activities of clostridial neurotoxins. The clostridial (including the  
CC Clostridium botulinum) neurotoxin substrate peptides are useful for  
CC detecting the presence of clostridial neurotoxins in a sample (e.g. food  
CC or an environmental sample). The present amino acid sequence represents a  
CC clostridial neurotoxin substrate peptide of the invention.  
XX  
SQ Sequence 116 AA;  
  
Query Match 100.0%; Score 39; DB 23; Length 116;  
Best Local Similarity 100.0%; Pred. No. 2.2;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 QIDRIMEX 8  
DB 86 QIDRIMEX 93  
  
RESULT 7  
AA015166  
ID AA015166 standard; peptide; 116 AA.  
XX  
XX AA015166;  
AC  
DT 02-SEP-2002 (first entry)  
XX  
XX Clostridial neurotoxin protease substrate peptide 5.  
DE  
XX  
XX Clostridial neurotoxin substrate; botulinum neurotoxin substrate; FRET;  
KM fluorescence resonant energy transfer assay; quenched-signal;  
KM clostridial neurotoxin detection; food.  
XX  
XX Unidentified.  
OS  
XX  
FH Key Location/Qualifiers  
FT Modified-site 1 /note= "S-fluoresceinyl-cysteine"  
FT Cleavage-site 89..90 /note= "The peptide is cleaved between these two  
FT residues by type E Clostridium botulinum neurotoxin"  
XX  
XX WO200225284-A2.

XX  
PD 28-MAR-2002.  
XX  
XX 25-SEP-2001; 2001WO-US30188.  
PF  
XX 25-SEP-2000; 2000US-235050P.  
PR  
XX (USME-) US MEDICAL RES INST INFECTIOUS DISEASES.  
PA Schmidt JI, Stafford RG;  
XX  
XX WPI: 2002-499829/53.  
XX  
XX Substrate useful in e.g. an assay for the protease activity of  
PT clostridial neurotoxin, comprises modified peptide or protein -  
PT  
XX  
XX Claim 28; Page 17; 48pp; English.  
XX  
XX The invention comprises clostridial neurotoxin substrate peptides which  
CC can serve as fluorescence resonant energy transfer assay (FRET) or  
CC quenched-signal substrates in assays for the proteolytic activities of  
CC clostridial neurotoxins. The invention further comprises Clostridium  
CC botulinum neurotoxin substrate peptides that can serve as immobilised  
CC substrates (i.e. bound to a solid phase) in assays for the proteolytic  
CC activities of clostridial neurotoxins. The clostridial (including the  
CC Clostridium botulinum) neurotoxin substrate peptides are useful for  
CC detecting the presence of clostridial neurotoxins in a sample (e.g. food  
CC or an environmental sample). The present amino acid sequence represents a  
CC clostridial neurotoxin substrate peptide of the invention.  
XX  
SQ Sequence 116 AA;  
  
Query Match 100.0%; Score 39; DB 23; Length 116;  
Best Local Similarity 100.0%; Pred. No. 2.2;  
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
QY 1 QIDRIMEX 8  
DB 86 QIDRIMEX 93  
  
RESULT 8  
AAU00255  
ID AAU00255 standard; Protein; 198 AA.  
XX  
XX AAU00255;  
AC  
DT 12-SEP-2001 (first entry)  
XX  
XX Synaptosomal-associated protein, SNAP25, C-terminal deletion 1-198.  
DE  
XX  
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KM toxin-inhibitory SNARE; botulinism; tetanus; SNARE-dependent exocytosis;  
KM synaptosomal-associated protein; mouse; mutant; mutein;  
KM N-ethylmaleimide-sensitive fusion protein;  
KM soluble NSF-attachment protein receptor.  
XX  
XX Mus sp.  
OS Synthetic.  
OS  
XX  
XX WO200118038-A2.  
PN  
XX  
XX 15-MAR-2001.  
PD  
XX  
XX 18-AUG-2000; 2000WO-GB03196.  
PF  
XX  
XX 20-AUG-1999; 99US-0149993.  
PR  
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
PA  
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
PI  
XX  
XX WPI: 2001-226739/23.

XX Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -  
 PS Example 1; Page - : 131pp; English.

XX The sequence represents the amino acid sequence of synaptosomal-  
 CC associated protein, SNAP25, C-terminal deletion 1-198, used in a new  
 CC method of treating a patient suffering from poisoning or at risk of  
 CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
 CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
 CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
 CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
 CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
 CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
 CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
 CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
 CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
 CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
 CC useful in the manufacture of a medicament for the treatment of a patient  
 CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
 CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
 CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
 CC polynucleotide encoding either of these SNARE polypeptides are useful in  
 CC the manufacture of medicament for the treatment of a patient in need of  
 CC inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis. The method of treatment is  
 CC relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC Note: The present sequence is not shown in the specification but is  
 CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).

SO Sequence 198 AA:  
 QY  
 Db 177 QIDRIMEX 184

Query Match 100.0%; Score 39; DB 22; Length 198;  
 Best Local Similarity 100.0%; Pred. No. 3.7;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

1 QIDRIMEX 8  
 |||||  
 177 QIDRIMEX 184

RESULT 9  
 AAU00263  
 ID AAU00263 standard; Protein: 199 AA.  
 AC AAU00263;  
 DT 12-SEP-2001 (first entry)  
 DE Synaptosomal-associated protein, SNAP25, mutant 1-199(R198T).  
 KW SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KW synaptosomal-associated protein; mouse; mutant; mutetin;  
 KW N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 OS Mus sp.  
 OS Synthetic.  
 FH Key Location/Qualifiers  
 FT Misc-difference 198 /note- "Wild-type Arg substituted by Thr"  
 PN WO200118038-A2.  
 PD 15-MAR-2001.  
 PF 18-AUG-2000; 2000MO-GB03196.  
 XX

PR 20-AUG-1999; 99US-0149993.  
 XX  
 PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 XX  
 PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
 XX  
 DR WPI; 2001-226739/23.  
 XX

PT Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -  
 PS Example 1; Page - : 131pp; English.

XX The sequence represents the amino acid sequence of synaptosomal-  
 CC associated protein, SNAP25, mutant 1-199(R198T), used in a new  
 CC method of treating a patient suffering from poisoning or at risk of  
 CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
 CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
 CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
 CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
 CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
 CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
 CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
 CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
 CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
 CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
 CC useful in the manufacture of a medicament for the treatment of a patient  
 CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
 CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
 CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
 CC polynucleotide encoding either of these SNARE polypeptides are useful in  
 CC the manufacture of medicament for the treatment of a patient in need of  
 CC inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis. The method of treatment is  
 CC relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC Note: The present sequence is not shown in the specification but is  
 CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).

SO Sequence 199 AA:  
 QY  
 Db 177 QIDRIMEX 184

Query Match 100.0%; Score 39; DB 22; Length 199;  
 Best Local Similarity 100.0%; Pred. No. 3.8;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

1 QIDRIMEX 8  
 |||||  
 177 QIDRIMEX 184

RESULT 10  
 AAU00264  
 ID AAU00264 standard; Protein: 200 AA.  
 AC AAU00264;  
 DT 12-SEP-2001 (first entry)  
 DE Synaptosomal-associated protein, SNAP25, mutant 1-200(R198T).  
 KW SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KW synaptosomal-associated protein; mouse; mutant; mutetin;  
 KW N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 OS Mus sp.  
 OS Synthetic.  
 FH Key Location/Qualifiers  
 FT Misc-difference 198 /note- "Wild-type Arg substituted by Thr"  
 PN  
 PD  
 PF  
 XX

```

XX  MO200118038-A2.
PN
XX
XX  15-MAR-2001.
PD
XX
XX  18-AUG-2000; 2000MO-GB03196.
PF
XX
XX  20-AUG-1999; 99US-0149993.
PR
XX
XX  (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
PA
XX  Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
PI
XX  WPI; 2001-226739/23.
PT
XX  Treating a patient suffering from poisoning or at risk of poisoning by
PT  a clostridial toxin, e.g. botulism, comprises administering a
PT  toxin-resistant or toxin-inhibitory SNARE -
PS
XX  Example 1; Page - : 131pp; English.
XX
XX  The sequence represents the amino acid sequence of synaptosomal-
CC  associated protein, SNAP25, mutant 1-200(R198T), used in a new
CC  method of treating a patient suffering from poisoning or at risk of
CC  poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
CC  (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
CC  to a cell of the patient, where the SNARE is resistant to proteolysis by
CC  the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
CC  toxin (toxin-inhibitory SNARE). The protein can be used in a method of
CC  treating a patient in need of inhibition of SNARE-dependent exocytosis
CC  from a cell capable of performing SNARE-dependent exocytosis, comprises
CC  supplying a fragment, variant, fusion or derivative of a SNARE or an
CC  inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
CC  inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
CC  useful in the manufacture of a medicament for the treatment of a patient
CC  suffering from poisoning or at risk of poisoning by clostridial toxin,
CC  e.g. from botulism or tetanus. The fragment, variant, fusion or
CC  derivative of a SNARE or of an inhibitory SNARE, or a recombinant
CC  polynucleotide encoding either of these SNARE polypeptides are useful in
CC  the manufacture of medicament for the treatment of a patient in need of
CC  inhibition of SNARE-dependent exocytosis from a cell capable of
CC  performing SNARE-dependent exocytosis. The method of treatment is
CC  relatively fast, thus alleviating the symptoms when most severe and
CC  taking the patient out of critical state.
CC  Note: The present sequence is not shown in the specification but is
CC  derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
CC
XX
XX  Sequence 200 AA:
SO
XX
XX  Query Match 100.0%; Score 39; DB 22; Length 200;
XX  Best Local Similarity 100.0%; Pred. No. 3.8;
XX  Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
OY 1 QIDRIMER 8
Db 177 QIDRIMER 184

```

```

XX  Mus sp.
OS Synthetic.
XX
XX  Key Location/Qualifiers
FT Misc-difference 198
FT /note="Wild-type Arg substituted by Thr"
PN
XX  MO200118038-A2.
PD
XX
XX  15-MAR-2001.
PF
XX
XX  18-AUG-2000; 2000MO-GB03196.
PR
XX
XX  20-AUG-1999; 99US-0149993.
PA
XX  (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
PI
XX  Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX  WPI; 2001-226739/23.
XX
XX  Treating a patient suffering from poisoning or at risk of poisoning by
PT  a clostridial toxin, e.g. botulism, comprises administering a
PT  toxin-resistant or toxin-inhibitory SNARE -
PS
XX  Example 1; Page - : 131pp; English.
XX
XX  The sequence represents the amino acid sequence of synaptosomal-
CC  associated protein, SNAP25, mutant 1-201(R198T), used in a new
CC  method of treating a patient suffering from poisoning or at risk of
CC  poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
CC  (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
CC  to a cell of the patient, where the SNARE is resistant to proteolysis by
CC  the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
CC  toxin (toxin-inhibitory SNARE). The protein can be used in a method of
CC  treating a patient in need of inhibition of SNARE-dependent exocytosis
CC  from a cell capable of performing SNARE-dependent exocytosis, comprises
CC  supplying a fragment, variant, fusion or derivative of a SNARE or an
CC  inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
CC  inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
CC  useful in the manufacture of a medicament for the treatment of a patient
CC  suffering from poisoning or at risk of poisoning by clostridial toxin,
CC  e.g. from botulism or tetanus. The fragment, variant, fusion or
CC  derivative of a SNARE or of an inhibitory SNARE, or a recombinant
CC  polynucleotide encoding either of these SNARE polypeptides are useful in
CC  the manufacture of medicament for the treatment of a patient in need of
CC  inhibition of SNARE-dependent exocytosis from a cell capable of
CC  performing SNARE-dependent exocytosis. The method of treatment is
CC  relatively fast, thus alleviating the symptoms when most severe and
CC  taking the patient out of critical state.
CC  Note: The present sequence is not shown in the specification but is
CC  derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
CC
XX
XX  Sequence 201 AA:
SO
XX
XX  Query Match 100.0%; Score 39; DB 22; Length 201;
XX  Best Local Similarity 100.0%; Pred. No. 3.8;
XX  Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
OY 1 QIDRIMER 8
Db 177 QIDRIMER 184

```

DE Synaptoosomal-associated protein, SNAP25, mutant 1-202(R198T).  
 XX  
 XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptoosomal-associated protein; mouse; mutant; mutein;  
 KM N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 XX  
 OS Mus sp.  
 OS Synthetic.  
 XX  
 FH Key Location/Qualifiers  
 FT MISC-difference 198  
 FT /note= "Wild-type Arg substituted by Thr"  
 XX  
 XX MO200118038-A2.  
 XX  
 PD 15-MAR-2001.  
 XX  
 PF 18-AUG-2000; 2000MO-GB03196.  
 XX  
 PF 20-AUG-1999; 99US-0149993.  
 XX  
 PR (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 XX  
 PI DOLLY JO, O'Sullivan GA, Mohammed N, Foran PG;  
 XX  
 DR WPI: 2001-226739/23.  
 XX  
 XX  
 PT Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -  
 XX  
 XX Example 1; Page - : 131pp; English.  
 PS  
 XX The sequence represents the amino acid sequence of synaptoosomal-  
 CC associated protein, SNAP25, mutant 1-202(R198T), used in a new  
 CC method of treating a patient suffering from poisoning or at risk of  
 CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
 CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
 CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
 CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
 CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
 CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
 CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
 CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
 CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
 CC inhibitory SNARE, or a recombinant polynucleotide encoding the SNARE is  
 CC useful in the manufacture of a medicament for the treatment of a patient  
 CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
 CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
 CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
 CC polynucleotide encoding either of these SNARE polypeptides are useful in  
 CC the manufacture of medicament for the treatment of a patient in need of  
 CC inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis. The method of treatment is  
 CC relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC Note: The present sequence is not shown in the specification but is  
 CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).  
 CC  
 XX  
 XX Sequence 202 AA;  
 SQ  
 Query Match 100.0%; Score 39; DB 22; Length 202;  
 Best Local Similarity 100.0%; Pred. No. 3.8;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 QIDRIMER 8  
 Db 177 QIDRIMER 184  
 RESULT 13

AAU02636  
 ID AAU02636 standard; Protein; 203 AA.  
 XX  
 XX AC AAU02636;  
 XX  
 XX 12-SEP-2001 (first entry)  
 DE Synaptoosomal-associated protein, SNAP25, mutant 1-203(R.98T).  
 XX  
 XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KM synaptoosomal-associated protein; mouse; mutant; mutein;  
 KM N-ethylmaleimide-sensitive fusion protein;  
 KM soluble NSF-attachment protein receptor.  
 XX  
 XX Mus sp.  
 OS Synthetic.  
 OS  
 FH Key Location/Qualifiers  
 FT MISC-difference 198  
 FT /note= "Wild-type Arg substituted by Thr"  
 XX  
 XX MO200118038-A2.  
 XX  
 PD 15-MAR-2001.  
 XX  
 PF 18-AUG-2000; 2000MO-GB03196.  
 XX  
 PF 20-AUG-1999; 99US-0149993.  
 XX  
 PR (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 XX  
 PI DOLLY JO, O'Sullivan GA, Mohammed N, Foran PG;  
 XX  
 DR WPI: 2001-226739/23.  
 XX  
 XX  
 PT Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -  
 XX  
 XX Example 1; Page - : 131pp; English.  
 PS  
 XX The sequence represents the amino acid sequence of synaptoosomal-  
 CC associated protein, SNAP25, mutant 1-203(R198T), used in a new  
 CC method of treating a patient suffering from poisoning or at risk of  
 CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
 CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
 CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
 CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
 CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
 CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
 CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
 CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
 CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
 CC inhibitory SNARE, or a recombinant polynucleotide encoding the SNARE is  
 CC useful in the manufacture of a medicament for the treatment of a patient  
 CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
 CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
 CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
 CC polynucleotide encoding either of these SNARE polypeptides are useful in  
 CC the manufacture of medicament for the treatment of a patient in need of  
 CC inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis. The method of treatment is  
 CC relatively fast, thus alleviating the symptoms when most severe and  
 CC taking the patient out of critical state.  
 CC Note: The present sequence is not shown in the specification but is  
 CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).  
 CC  
 XX  
 XX Sequence 203 AA;  
 SQ  
 Query Match 100.0%; Score 39; DB 22; Length 203;  
 Best Local Similarity 100.0%; Pred. No. 3.8;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 QIDRIMEX 8  
 ID |||||||  
 Db 177 QIDRIMEX 184

RESULT 14  
 AAM30103  
 ID AAM30103 standard; peptide; 206 AA.

XX AAM30103;

XX 06-APR-1998 (first entry)

XX Synaptosomal associated protein.

XX Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;  
 KM excitation-secretory uncoupling peptide; catecholamine secretion;  
 KM bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
 KM synaptosomal associated protein; SNAP-25.

XX Homo sapiens.

XX MO9734620-A1.

XX 25-SEP-1997.

XX 18-MAR-1997; 97MO-US04393.

XX 18-MAR-1996; 96US-0013599.

XX (REGC) UNIV CALIFORNIA.

XX Montal M;

XX WPI; 1997-479986/44.

XX Excitation-secretory uncoupling peptide(s) for inhibiting  
 PT neuro-transmitter release - used particularly for treating muscle  
 PT spasticity, and for delivering drugs specifically to neural cells

XX Disclosure: Page 27-28; 61pp; English.

XX This sequence represents the human 25 kD synaptosomal associated protein  
 CC (SNAP-25), which is an inhibitory agent of the invention. The agents of  
 CC the invention inhibit secretion of neurotransmitter from neuronal cells  
 CC and is an excitation-secretory uncoupling peptide (1) of at least 20  
 CC amino acids (aa) all of which correspond substantially to any one of  
 CC AAM30097-W30102, or more generally any (1) that inhibits 50% of  
 CC catecholamine secretion from bovine chromaffin cells at a concentration  
 CC of 10 microm, especially 0.25 microm, or less. (1) are used, as a  
 CC replacement for Clostridium toxin, to inhibit release of  
 CC neurotransmitters from synaptic vesicles, specifically for reducing  
 CC muscle spasticity. Also (1) may be labelled to allow in vivo imaging of  
 CC intracellular distribution of (1). Compounds for delivering the drug to  
 CC neural cells provide targeted drug delivery, e.g. of substance P to  
 CC brain tumours for induction of apoptosis. Unlike the neurotoxins, (1) are  
 CC not toxic or immunogenic and are more readily available. Their  
 CC therapeutic effect lasts for several days or weeks, so lower doses or  
 CC less frequent treatments are required.

XX Sequence 206 AA;

Query Match 100.0%; Score 39; DB 18; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 3.9;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 QIDRIMEX 8  
 ID |||||||  
 Db 177 QIDRIMEX 184

RESULT 15

AAM79198  
 ID AAM79198 standard; Protein; 206 AA.

XX AAM79198;

XX 25-NOV-1998 (first entry)

XX Mouse SNAP-25 polypeptide.

XX Hrs-2 polypeptide; AMP-preferring nucleotidase; SNAP-25; vesicle docking;  
 KM calcium-regulated secretion; secretory vesicle; secretory process; brain;  
 KM neurotransmitter release; presynaptic membrane; CNS disorder; depression;  
 KM Parkinson's disease; endocrine system; hormonal imbalance; cell division;  
 KM thought disorder; schizophrenia; degenerative disorder; anaesthesia; rat;  
 KM immune system; antigen processing; immunomodulator; viral processing;  
 KM central nervous system; vesicular release; vesicular disorder; human;  
 KM anti-tumour application; membrane trafficking regulation; mouse.

XX Mus sp.

XX MO9838210-A2.

XX 03-SEP-1998.

XX 26-FEB-1998; 98MO-US03789.

XX 26-FEB-1997; 97US-0039159.

XX (STRD) UNIV LELAND STANFORD JUNIOR.

XX Bean AJ, Scheller RH;

XX WPI; 1998-481140/41.

XX N-PsDB; AAV57558.

XX New isolated Hrs-2 nucleotidase - used in assays to identify  
 PT compounds capable of modulating calcium-regulatory secretion of  
 PT secretory vesicles, such as in neurotransmitter release  
 PS Claim 16; Pages 42-44; 55pp; English.

XX This represents a mouse SNAP-25 polypeptide, a component of the protein  
 CC polypeptides thought to underlie vesicle docking and fusion. The  
 CC invention provides rat and human Hrs-2 polypeptides which are ATP-  
 CC preferring nucleotidase that associate with SNAP-25. For identifying a  
 CC compound capable of modulating calcium-regulated secretion of secretory  
 CC vesicles, a SNAP-25 polypeptide can be contacted with a Hrs-2  
 CC polypeptide. In the presence and absence of a test compound. The effect  
 CC of the test compound on the extent of binding between the SNAP-25 and  
 CC Hrs-2 polypeptides are measured and a compound is identified as effective  
 CC if its measured effect on the extent of binding is above a threshold  
 CC level. The products can be used for identifying drugs capable of  
 CC affecting secretory processes, such as neurotransmitter release at the  
 CC active zones of presynaptic membranes. Such drugs can be used for  
 CC treating disorders or conditions of the central nervous system by  
 CC selectively enhancing or inhibiting vesicular release in specific areas  
 CC of the brain, including affective disorders (e.g. depression), disorders  
 CC of thought (e.g. schizophrenia) and degenerative disorders (Parkinson's  
 CC disease), as well as applications such as anaesthesia. The drugs can  
 CC also be used therapeutically in other systems such as the endocrine  
 CC system for treatment of hormonal imbalances, the immune system for  
 CC intervention in antigen processing, secreted immunomodulators, and viral  
 CC processing, as well as anti-tumour applications, such as regulation of  
 CC membrane trafficking during rapid cell division.

XX Sequence 206 AA;

Query Match 100.0%; Score 39; DB 19; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 3.9;  
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 QIDRIMEX 8  
 ID |||||||



Tue Dec 3 08:36:12 2002

pct-us02-27145-8.rag

Page 9

Db 177 QIDRIMEK 184

Search completed: November 19, 2002, 17:34:27  
Job time : 4.35135 secs

---

**THIS PAGE BLANK (USPTO)**

GenCore version 5.1.3  
Copyright (c) 1993 - 2002 Compugen Ltd.

OM protein - protein search, using sw model

Run on: November 19, 2002, 17:35:04 : Search time 22.3125 Seconds  
(without alignments)  
101.524 Million cell updates/sec

Title: PCT-US02-27145-2\_COPY\_187\_203  
Perfect score: 83  
Sequence: 1 SNKTRIDEANORATKML 17

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database :

A\_Geneseq\_101002:.\*  
1: /SID52/gcgdata/geneseq/geneseq-emb1/AA1980.DAT:\*  
2: /SID52/gcgdata/geneseq/geneseq-emb1/AA1981.DAT:\*  
3: /SID52/gcgdata/geneseq/geneseq-emb1/AA1982.DAT:\*  
4: /SID52/gcgdata/geneseq/geneseq-emb1/AA1983.DAT:\*  
5: /SID52/gcgdata/geneseq/geneseq-emb1/AA1984.DAT:\*  
6: /SID52/gcgdata/geneseq/geneseq-emb1/AA1985.DAT:\*  
7: /SID52/gcgdata/geneseq/geneseq-emb1/AA1986.DAT:\*  
8: /SID52/gcgdata/geneseq/geneseq-emb1/AA1987.DAT:\*  
9: /SID52/gcgdata/geneseq/geneseq-emb1/AA1988.DAT:\*  
10: /SID52/gcgdata/geneseq/geneseq-emb1/AA1989.DAT:\*  
11: /SID52/gcgdata/geneseq/geneseq-emb1/AA1990.DAT:\*  
12: /SID52/gcgdata/geneseq/geneseq-emb1/AA1991.DAT:\*  
13: /SID52/gcgdata/geneseq/geneseq-emb1/AA1992.DAT:\*  
14: /SID52/gcgdata/geneseq/geneseq-emb1/AA1993.DAT:\*  
15: /SID52/gcgdata/geneseq/geneseq-emb1/AA1994.DAT:\*  
16: /SID52/gcgdata/geneseq/geneseq-emb1/AA1995.DAT:\*  
17: /SID52/gcgdata/geneseq/geneseq-emb1/AA1996.DAT:\*  
18: /SID52/gcgdata/geneseq/geneseq-emb1/AA1997.DAT:\*  
19: /SID52/gcgdata/geneseq/geneseq-emb1/AA1998.DAT:\*  
20: /SID52/gcgdata/geneseq/geneseq-emb1/AA1999.DAT:\*  
21: /SID52/gcgdata/geneseq/geneseq-emb1/AA2000.DAT:\*  
22: /SID52/gcgdata/geneseq/geneseq-emb1/AA2001.DAT:\*  
23: /SID52/gcgdata/geneseq/geneseq-emb1/AA2002.DAT:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	83	100.0	17	20	AAV44021	Amino acids 187-20
2	83	100.0	17	20	AAV44057	Human SNAP-25 (amin
3	83	100.0	17	23	ABG69065	Human polypeptide
4	83	100.0	19	22	AA15366	Human SNAP-25 N-te
5	83	100.0	20	18	AAW30100	Neurotransmitter s
6	83	100.0	26	18	AAW30099	Neurotransmitter s
7	83	100.0	37	18	AAW30097	Neurotransmitter s
8	83	100.0	70	17	AAW68823	SNAP-25 residues 1
9	83	100.0	86	22	AA15584	Human SNAP-25 N-te
10	83	100.0	116	23	AA015165	Clostridial neurot

11	83	100.0	206	18	AAW30103	;;synaptosomal assoc
12	83	100.0	206	19	AAW79198	house SNAP-25 poly
13	83	100.0	206	19	AAW43426	house synaptosomal
14	83	100.0	206	22	AAU00246	;;synaptosomal-assoc
15	83	100.0	206	22	AAU00252	;;SNARE homologue, s
16	83	100.0	206	22	AAU00253	;;SNARE homologue, s
17	80	96.4	17	20	AAW44038	;;human SNAP25 (amin
18	80	96.4	17	20	AAW44063	;;human SNAP25 (amin
19	79	95.2	16	20	AAW44069	;;human SNAP25 (amin
20	79	95.2	17	20	AAW44047	;;human SNAP25 (amin
21	79	95.2	17	20	AAW44050	;;human SNAP25 (amin
22	79	95.2	17	20	AAW44052	;;human SNAP25 (amin
23	79	95.2	17	20	AAW44059	;;human SNAP25 (amin
24	79	95.2	206	22	AAU02640	;;synaptosomal-assoc
25	78	94.0	17	20	AAW44039	;;human SNAP25 (amin
26	78	94.0	17	20	AAW44045	;;human SNAP25 (amin
27	78	94.0	17	20	AAW44049	;;human SNAP25 (amin
28	78	94.0	17	20	AAW44062	;;human SNAP25 (amin
29	78	94.0	17	20	AAW44070	;;human SNAP25 (amin
30	77	92.8	17	20	AAW44022	;;human SNAP25 (amin
31	77	92.8	17	20	AAW44040	;;human SNAP25 (amin
32	77	92.8	17	20	AAW44044	;;human SNAP25 (amin
33	77	92.8	17	20	AAW44046	;;human SNAP25 (amin
34	77	92.8	17	20	AAW44048	;;human SNAP25 (amin
35	77	92.8	17	20	AAW44051	;;human SNAP25 (amin
36	77	92.8	17	20	AAW44053	;;human SNAP25 (amin
37	77	92.8	17	20	AAW44054	;;human SNAP25 (amin
38	77	92.8	17	20	AAW44056	;;human SNAP25 (amin
39	77	92.8	17	20	AAW44064	;;human SNAP25 (amin
40	77	92.8	17	20	AAW44065	;;human SNAP25 (amin
41	77	92.8	17	20	AAW44066	;;human SNAP25 (amin
42	77	92.8	24	23	AA015162	;;Clostridial neurot
43	77	92.8	116	23	AA015166	;;Clostridial neurot
44	77	92.8	203	22	AAU02636	;;synaptosomal-assoc
45	77	92.8	206	22	AAU00259	;;synaptosomal-assoc

#### ALIGNMENTS

RESULT 1	
AAV44021	
ID	AAV44021 standard; peptide; 17 AA.
XX	
AC	AAV44021;
XX	
DT	18-JAN-2000 (first entry)
XX	
DE	Amino acids 187-203 of human SNAP25.
XX	
KW	Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;
KW	fluorescamine; detection; human; synaptosomal protein; SNAP25;
KW	hydrolysis; amino group.
XX	
OS	Homo sapiens.
XX	
PN	US5965699-A.
XX	
PD	12-OCT-1999.
XX	
PF	06-NOV-1996; 96US-0743894.
XX	
PR	06-NOV-1996; 96US-0743894.
XX	
PA	(USSA ) US SEC OF ARMY.
XX	
PI	Bostian KA, Schmidt JF;
XX	
DR	WPI; 1999-579939/49.
XX	
PT	Quantitation of type A botulinum toxin -
XX	
PS	Claim 1; Column 4; 28pp; English.

```

XX CC The invention relates to an enzymatic assay for the quantitation of
CC CC type A botulinum toxin, by determining the proteolytic activity of
CC CC botulinum neurotoxin type A using fluorescamine detection. The method
CC CC comprises adding an analogue (e.g. AA44022-Y44076) of this peptide
CC CC (which represents amino acids 187-203 of the human synaptosomal protein
CC CC SNAP25) to a sample containing the botulinum toxin A so that hydrolysis
CC CC of the peptide is initiated, then stopping hydrolysis of the peptide at
CC CC different time points; and measuring the amount of hydrolysis at each
CC CC time point by combining with a label capable of detecting free amino
CC CC groups resulting from the hydrolysis. The amount of botulinum toxin A
CC CC present in the sample is determined by comparing measurements with the
CC CC amount of label produced from a known concentration of toxin measured
CC CC under similar conditions. The method is useful for the quantitation of
CC CC type A botulinum toxin.
XX

SQ Sequence 17 AA:

Query Match 100.0%; Score 83; DB 20; Length 17;
Best Local Similarity 100.0%; Pred. No. 9.4e-08;
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 SNKTRIDEANORATKML 17
DB 1 SNKTRIDEANORATKML 17
|||||
|||||

RESULT 2
AA444057
ID AA444057 standard; peptide; 17 AA.
XX
XX AA444057;
XX
XX 18-JAN-2000 (first entry)
XX
XX Human SNAP25 (amino acids 187-203) analogue #36.
XX
XX Enzymatic assay; quantitation; type A botulinum neurotoxin; proteolysis;
XX KM fluorescamine; detection; human; synaptosomal protein; SNAP25;
XX KM hydrolysis; amino group.
XX
XX Homo sapiens.
XX OS Synthetic.
XX
XX US5965699-A.
XX PN 12-OCT-1999.
XX
XX 06-NOV-1996; 96US-0743894.
XX PF
XX 06-NOV-1996; 96US-0743894.
XX PR
XX 06-NOV-1996; 96US-0743894.
XX
XX (USSA ) US SEC OF ARMY.
XX PA
XX Bostlian KA, Schmidt JJ;
XX PI
XX WPI; 1999-579939/49.
XX
XX Quantitation of type A botulinum toxin -
XX
XX Disclosure; Column 9; 28pp; English.
XX
XX The invention relates to an enzymatic assay for the quantitation of
XX CC type A botulinum toxin, by determining the proteolytic activity of
XX CC botulinum neurotoxin type A using fluorescamine detection. Botulinum
XX CC toxin A has been shown to cleave the synaptosomal neurotransmitter
XX CC peptide SNAP25 between residues 197-198. The method comprises adding
XX CC an analogue (e.g. AA44022-Y44076) of the SNAP25 peptide (AA44021,
XX CC amino acids 187-203 of human SNAP25) to a sample containing the
XX CC botulinum toxin A so that hydrolysis of the peptide is initiated, then
XX CC stopping hydrolysis of the peptide at different time points; and
XX CC measuring the amount of hydrolysis at each time point by combining with a
XX CC label capable of detecting free amino groups resulting from the
XX

```

```

CC CC hydrolysis. The amount of botulinum toxin A present in the sample is
CC CC determined by comparing measurements with the amount of label produced
CC CC from a known concentration of toxin measured under similar conditions.
CC CC The method is useful for the quantitation of type A botulinum toxin.
XX

SQ Sequence 17 AA:

Query Match 100.0%; Score 83; DB 20; Length 17;
Best Local Similarity 100.0%; Pred. No. 9.4e-08;
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 SNKTRIDEANORATKML 17
DB 1 SNKTRIDEANORATKML 17
|||||
|||||

RESULT 3
ABG69065
ID ABG69065 standard; Peptide; 17 AA.
XX
XX ABG69065;
XX
XX 07-OCT-2002 (first entry)
XX
XX Human polypeptide C-terminal fragment.
XX
XX Botulinum neurotoxin light chain; BoNT LC; botulism; dystonia; pain;
XX KM spasticity; ocular motility; facial dyskinesia; stiff-person syndrome;
XX KM bladder dysfunction; segmental myoclonus; hyperkinetic disorder; human;
XX KM cosmetic treatment; facial wrinkle; cerebral palsy; analgesic; relaxant;
XX KM lower motor neuron hyperactivity; autonomic nerve function; muscular;
XX KM immunostimulant; antibacterial.
XX
XX Homo sapiens.
XX OS
XX
XX WO200236758-A2.
XX PN 10-MAY-2002.
XX
XX 06-NOV-2001; 2001WO-US47230.
XX PF
XX 06-NOV-2000; 2000US-246774P.
XX PR 20-JUL-2001; 2001US-0910186.
XX PR 09-AUG-2001; 2001US-311966P.
XX
XX (USSA ) US ARMY MEDICAL RES & MATERIAL COMMAND.
XX PA
XX Smith LA, Jensen M;
XX PI
XX WPI; 2002-575192/61.
XX
XX Novel nucleic acid molecule encoding botulinum neurotoxin light chain
XX PT serotype A, useful for producing the neurotoxin for vaccination against
XX PT botulism, comprises sequence expressible in host other than Clostridium
XX
XX Example 25; Page 62; 16pp; English.
XX
XX The invention relates to a nucleic acid molecule encoding a botulinum
XX CC neurotoxin light chain (BoNT LC) serotype A, where the DNA has a sequence
XX CC that is expressible in a host organism other than Clostridium, or has a
XX CC total A+T content that is less than about 70% The BoNT LC protein is
XX CC useful in vaccination against botulism, for eliciting protective immunity
XX CC in a mammal, for treating dystonias, spasticity, pain, ocular motility,
XX CC facial dyskinesias, stiff-person syndrome, bladder dysfunction, segmental
XX CC myoclonus, hyperkinetic disorders, cosmetic treatment of facial wrinkles,
XX CC conditions characterised by hyperactivity of the lower motor neuron, and
XX CC to control autonomic nerve function or lipoe-walking due to stiff
XX CC muscles common in children with cerebral palsy. The sequences are also
XX CC useful for screening for botulinum neurotoxin inhibitors. This sequence
XX CC represents a human polypeptide C-terminal fragment, used in the scope of
XX CC the invention.
XX

```

SO Sequence 17 AA;  
Query Match 100.0%; Score 83; DB 23; Length 17;  
Best Local Similarity 100.0%; Pred. No. 9.4e-08;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 SNKTRIDEANORATKML 17  
1 SNKTRIDEANORATKML 17  
DB 1 SNKTRIDEANORATKML 17  
RESULT 4  
AAB15586  
ID AAB15586 standard; peptide: 19 AA.  
XX  
AC AAB15586;  
XX  
DT 02-MAR-2001 (first entry)  
XX  
DE Human SNAP-25 N-terminal peptide #6.  
XX  
DE Dermatological; neuroprotective; relaxant; neuronal exocytosis modulator;  
KW SNAP-25; synaptosomal associated protein 25; facial wrinkle; asymmetry;  
KW neurodegenerative disorder.  
XX  
OS Homo sapiens.  
XX  
PN WO200064932-A1.  
XX  
PD 02-NOV-2000.  
XX  
PE 18-FEB-2000; 2000WO-ES00058.  
XX  
PR 23-APR-1999; 99ES-0000844.  
XX  
PA (LIPD-) LIPOTEC SA.  
XX  
PI Blanes Mira MC, Llobregat Hernandez MM, Gil Tebar AI;  
PI Fernandez Ballesster GJ, Planell Cases RM, Ferrer Montiel AV;  
PI Viniegra Bover S, Gutierrez Perez LM, Carbonell Castell T;  
PI Perez Paya E;  
XX  
DR WPI: 2001-007091/01.  
XX  
PT New peptides containing amino acid sequences from known proteins for  
PT treatment of neurological disorders  
XX  
PS Claim 17; Page 34; 40pp; Spanish.  
XX  
CC The invention relates to new peptides comprising 3-30 contiguous amino  
CC acids from the N-terminus of the protein SNAP-25  
CC (synaptosomal-associated protein 25). The peptides AAB15581-B15586  
CC represent examples of the peptides of the invention. The peptides have  
CC neuronal exocytosis inhibitory activity and are used for treatment of  
CC facial wrinkles and asymmetry and pathological neuronal  
CC exocytosis-mediated pathological disorders and alterations manifested  
CC e.g. by spasms and neurological and neurodegenerative disorders.  
XX  
SQ Sequence 19 AA;  
Query Match 100.0%; Score 83; DB 22; Length 19;  
Best Local Similarity 100.0%; Pred. No. 1.1e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 SNKTRIDEANORATKML 17  
1 SNKTRIDEANORATKML 17  
DB 3 SNKTRIDEANORATKML 19  
RESULT 5  
AAW30100  
ID AAW30100 standard; peptide: 20 AA.  
XX

AC AAW30100;  
XX  
DT 06-APR-1998 (first entry)  
XX  
DE Neurotransmitter secretion inhibitor #4.  
XX  
KW Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;  
KW excitation-secretory uncoupling peptide; catecholamine secretion;  
KW bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
KW synaptosomal associated protein; SNAP-25.  
XX  
OS Homo sapiens.  
XX  
PN WO9734620-A1.  
XX  
PD 25-SEP-1997.  
XX  
PE 18-MAR-1997; 97WO-US04393.  
XX  
PR 18-MAR-1996; 96US-0013599.  
XX  
PA (RBC ) UNIV CALIFORNIA.  
PI Montal M;  
XX  
DR WPI: 1997-479986/44.  
XX  
PT Excitation-secretory uncoupling peptide(s) for inhibiting  
PT neuro-transmitter release - used particularly for treating muscle  
PT spasticity, and for delivering drugs specifically to mural cells  
XX  
PS Claim 14; Page 32; 61pp; English.  
XX  
CC This sequence corresponds to residues 187-206 of the human 25 kD  
CC synaptosomal associated protein (SNAP-25), and is a inhibitory agent of  
CC the invention. The agents of the invention inhibit secretion of  
CC neurotransmitter from neuronal cells and is an exciton-secretory  
CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which  
CC correspond substantially to any one of AAW30097-W30102, or more  
CC generally any (I) that inhibits 50% of catecholamine secretion from  
CC bovine chromaffin cells at a concentration of 10 microM, especially 0.25  
CC microM, or less. (I) are used, as a replacement for Clostridium toxin, to  
CC inhibit release of neurotransmitters from synaptic vesicles, specifically  
CC for reducing muscle spasticity. Also (I) may be labelled to allow in  
CC vivo imaging of intracellular distribution of (I). Compounds for  
CC delivering the drug to neural cells provide targeted drug delivery, e.g.  
CC of substance P to brain tumours for induction of apoptosis. Unlike the  
CC neurotoxins, (I) are not toxic or immunogenic and are more readily  
CC available. Their therapeutic effect lasts for several days or weeks, so  
CC lower doses or less frequent treatments are required.  
XX  
SQ Sequence 20 AA;  
Query Match 100.0%; Score 83; DB 18; Length 20;  
Best Local Similarity 100.0%; Pred. No. 1.1e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 SNKTRIDEANORATKML 17  
1 SNKTRIDEANORATKML 17  
DB 1 SNKTRIDEANORATKML 17  
RESULT 6  
AAW30099  
ID AAW30099 standard; peptide: 26 AA.  
XX  
AC AAW30099;  
XX  
DT 06-APR-1998 (first entry)  
XX  
DE Neurotransmitter secretion inhibitor #3.  
KW Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;

KW excitation-secretory uncoupling peptide; catecholamine secretion;  
KW bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
KW synaptosomal associated protein; SNAP-25.  
XX  
OS Homo sapiens.  
XX  
PN WO9734620-A1.  
XX  
PD 25-SEP-1997.  
XX  
PF 18-MAR-1997; 97WO-US04393.  
XX  
PR 18-MAR-1996; 96US-0013599.  
XX  
PA (RECC ) UNIV CALIFORNIA.  
XX  
PI Montal M;  
XX  
PS WPI; 1997-479986/44.  
DR  
XX  
XX  
PT Excitation-secretory uncoupling peptide(s) for inhibiting  
PT neuro:transmitter release - used particularly for treating muscle  
PT spasticity, and for delivering drugs specifically to neural cells  
XX  
XX  
PS Claim 13; Page 31; 61pp; English.  
XX  
XX  
XX This sequence corresponds to residues 181-206 of the human 25 kD  
CC synaptosomal associated protein (SNAP-25), and is a inhibitory agent of  
CC the invention. The agents of the invention inhibit secretion of  
CC neurotransmitter from neuronal cells and is an excitation-secretory  
CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which  
CC correspond substantially to any one of AAW30097-W30102, or more  
CC generally any (I) that inhibits 50% of catecholamine secretion from  
CC bovine chromaffin cells at a concentration of 10 microm, especially 0.25  
CC microm, or less. (I) are used, as a replacement for Clostridium toxin, to  
CC inhibit release of neurotransmitters from synaptic vesicles, specifically  
CC for reducing muscle spasticity. Also (I) may be labelled to allow in  
CC vivo imaging of intracellular distribution of (I). Compounds for  
CC delivering the drug to neural cells provide targeted drug delivery, e.g.  
CC of substance P to brain tumours for induction of apoptosis. Unlike the  
CC neurotoxins, (I) are not toxic or immunogenic and are more readily  
CC available. Their therapeutic effect lasts for several days or weeks, so  
CC lower doses or less frequent treatments are required.  
XX  
SQ Sequence 26 AA:  
Query Match 100.0%; Score 83; DB 18; Length 26;  
Best Local Similarity 100.0%; Pred. No. 1.5e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 SNKTRIDEANQRATKML 17  
DB 7 SNKTRIDEANQRATKML 23  
|||||  
RESULT 7  
AAW30097 standard; peptide: 37 AA.  
XX  
AC AAW30097;  
XX  
DT 06-APR-1998 (first entry)  
XX  
DE Neurotransmitter secretion inhibitor #1.  
XX  
XX Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;  
KW excitation-secretory uncoupling peptide; catecholamine secretion;  
KW bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
KW synaptosomal associated protein; SNAP-25.  
XX  
XX Homo sapiens.  
XX  
PM WO9734620-A1.

XX  
XX 25-SEP-1997.  
XX  
XX 18-MAR-1997; 97WO-US04393.  
XX  
XX 18-MAR-1996; 96US-0013599.  
XX  
XX (RECC ) UNIV CALIFORNIA.  
XX  
XX Montal M;  
XX  
XX WPI; 1997-479986/44.  
DR  
XX  
XX  
PT Excitation-secretory uncoupling peptide(s) for inhibiting  
PT neuro:transmitter release - used particularly for treating muscle  
PT spasticity, and for delivering drugs specifically to neural cells  
XX  
XX  
PS Claim 1; Page 30; 61pp; English.  
XX  
XX  
XX This sequence corresponds to residues 170-206 of the human 25 kD  
CC synaptosomal associated protein (SNAP-25), and is a inhibitory agent of  
CC the invention. The agents of the invention inhibit secretion of  
CC neurotransmitter from neuronal cells and is an excitation-secretory  
CC uncoupling peptide (I) of at least 20 amino acids (aa) all of which  
CC correspond substantially to any one of AAW30097-W30102, or more  
CC generally any (I) that inhibits 50% of catecholamine secretion from  
CC bovine chromaffin cells at a concentration of 10 microm, especially 0.25  
CC microm, or less. (I) are used, as a replacement for Clostridium toxin, to  
CC inhibit release of neurotransmitters from synaptic vesicles, specifically  
CC for reducing muscle spasticity. Also (I) may be labelled to allow in  
CC vivo imaging of intracellular distribution of (I). Compounds for  
CC delivering the drug to neural cells provide targeted drug delivery, e.g.  
CC of substance P to brain tumours for induction of apoptosis. Unlike the  
CC neurotoxins, (I) are not toxic or immunogenic and are more readily  
CC available. Their therapeutic effect lasts for several days or weeks, so  
CC lower doses or less frequent treatments are required.  
XX  
SQ Sequence 37 AA:  
Query Match 100.0%; Score 83; DB 18; Length 37;  
Best Local Similarity 100.0%; Pred. No. 2.3e-07;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 SNKTRIDEANQRATKML 17  
DB 18 SNKTRIDEANQRATKML 34  
|||||  
RESULT 8  
AAR86823 standard; Peptide: 70 AA.  
XX  
AC AAR86823;  
XX  
DT 15-AUG-1996 (first entry)  
XX  
DE SNAP-25 residues 137-206.  
XX  
XX VAMP; vesicle-associated membrane protein; SNAP-25; syntaxin;  
KW neurotransmitter; neurotoxin; botulinum; botulism; cleavage;  
KW substrate; antibody; detection; assay.  
XX  
XX Synthetic.  
XX  
XX WO9533850-A1.  
XX  
XX 14-DEC-1995.  
XX  
XX 02-JUN-1995; 95WO-GB01279.  
XX  
XX 03-JUN-1994; 94GB-0011138.  
XX  
XX (CAMR-) CAMR CENT APPLIED MICROBIOLOGY & RES.

PA (MICR-) MICROBIOLOGICAL RES AUTHORITY.  
 XX  
 PI Hallis B, James BAF, Quinn CP, Shone CC;  
 XX  
 DR WPI: 1996-040249/04.  
 XX  
 PT Assay for botulinum or tetanus toxin - by combining test cpd. with  
 PT substrate which is cleaved by the toxin, and antibody specific for  
 PT the cleaved but not uncleaved substrate  
 XX  
 PS Example 4: Page 19; 48pp; English.  
 XX  
 CC The botulinum neurotoxins possess highly specific zinc-endopeptidase  
 CC activities within their light sub-units. Depending on the neurotoxin  
 CC type these act to cleave small proteins within the nerve cell which are  
 CC involved in neurotransmitter release. Antibodies are used in assays  
 CC which detect cleaved but not uncleaved substrate. Assays for botulinum  
 CC types A and E use the present sequence as a substrate. The sequence is  
 CC SNAP-25 protein, residues 137-206.  
 XX  
 SQ Sequence 70 AA:  
 XX  
 Query Match 100.0%; Score 83; DB 17; Length 70;  
 Best Local Similarity 100.0%; Pred. No. 4.6e-07;  
 Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 OY 1 SNKTRIDEANQRATKML 17  
 Db 51 SNKTRIDEANQRATKML 67  
 |||||||||||||||  
 RESULT 9  
 AAB15584  
 ID AAB15584 standard; peptide: 86 AA.  
 XX  
 AC AAB15584:  
 XX  
 DT 02-MAR-2001 (first entry)  
 XX  
 DE Human SNAP-25 N-terminal peptide #4.  
 XX  
 KW Dermatological; neuroprotective; relaxant; neuronal exocytosis modulator;  
 KW SNAP-25; synaptosomal-associated protein 25; facial wrinkle; asymmetry;  
 KW neurodegenerative disorder.  
 XX  
 OS Homo sapiens.  
 XX  
 PN MO200064932-A1.  
 XX  
 PD 02-NOV-2000.  
 XX  
 PF 18-FEB-2000; 2000WO-ES00058.  
 XX  
 PR 23-APR-1999; 99ES-0000844.  
 XX  
 PA (LIP0-) LIPOTEC SA.  
 XX  
 PI Blanes Mira MC, Llobregat Hernandez MM, Gil Tebar AI;  
 PI Fernandez Ballester GJ, Planell Cases RM, Ferrer Montiel AV;  
 PI Vinagre Bover S, Gutierrez Perez LM, Carbonell Castell T;  
 PI Perez Paya E;  
 XX  
 DR WPI: 2001-007091/01.  
 XX  
 PT New peptides containing amino acid sequences from known proteins for  
 PT treatment of neurological disorders  
 XX  
 PS Claim 16; Page 32-33; 40pp; Spanish.  
 XX  
 CC The invention relates to new peptides comprising 3-30 contiguous amino  
 CC acids from the N-terminus of the protein SNAP-25  
 CC (synaptosomal-associated protein 25). The peptides AAB15581-B15586  
 CC represent examples of the peptides of the invention. The peptides have

CC neuronal exocytosis inhibitory activity and are used for treatment of  
 CC facial wrinkles and asymmetry and pathological neuronal  
 CC exocytosis-mediated pathological disorders and alterations manifested  
 CC e.g. by spasms and neurological and neurodegenerative disorders.  
 XX  
 SQ Sequence 86 AA:  
 XX  
 Query Match 100.0%; Score 83; DB 22; Length 86;  
 Best Local Similarity 100.0%; Pred. No. 5.8e-07;  
 Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 OY 1 SNKTRIDEANQRATKML 17  
 Db 67 SNKTRIDEANQRATKML 83  
 |||||||||||||||  
 RESULT 10  
 AAO15165  
 ID AAO15165 standard; peptide: 116 AA.  
 XX  
 AC AAO15165:  
 XX  
 DT 02-SEP-2002 (first entry)  
 XX  
 DE Clostridial neurotoxin protease substrate peptide 4.  
 XX  
 KW Clostridial neurotoxin substrate; botulinum neurotoxin substrate; FRRT;  
 KW fluorescence resonant energy transfer assay; quenched-signal;  
 KW clostridial neurotoxin detection; food.  
 XX  
 OS Unidentified.  
 XX  
 FH Key Location/Qualifiers  
 FT Modified-site 1 /note= "S-fluoresceinyl-cysteine"  
 FT Cleavage-site 89..90 /note= "The peptide is cleaved between these two  
 FT residues by type E Clostridium botulinum neurotoxin"  
 FT Cleavage-site 106..107 /note= "The peptide is cleaved between these two  
 FT residues by type A Clostridium botulinum neurotoxin"  
 XX  
 PN MO200225284-A2.  
 XX  
 PD 28-MAR-2002.  
 XX  
 PF 25-SEP-2001; 2001WO-US30188.  
 XX  
 PR 25-SEP-2000; 2000US-235050P.  
 XX  
 PA (USME-) US MEDICAL RES INST INFECTIOUS DISEASES.  
 XX  
 PI Schmidt JJ, Stafford RG;  
 XX  
 DR WPI: 2002-499829/53.  
 XX  
 PT Substrate useful in e.g. an assay for the protease activity of  
 PT clostridial neurotoxin, comprises modified peptide or protein  
 XX  
 PS Claim 22; Page 16; 48pp; English.  
 XX  
 CC The invention comprises clostridial neurotoxin substrate peptides which  
 CC can serve as fluorescence resonant energy transfer assay (FRRT) or  
 CC quenched-signal substrates in assays for the proteolytic activities of  
 CC clostridial neurotoxins. The invention further comprises Clostridium  
 CC botulinum neurotoxin substrate peptides that can serve as immobilised  
 CC substrates (i.e. bound to a solid phase) in assays for the proteolytic  
 CC activities of clostridial neurotoxins. The clostridial (including the  
 CC Clostridium botulinum) neurotoxin substrate peptides are useful for  
 CC detecting the presence of clostridial neurotoxins in a sample (e.g. food  
 CC or an environmental sample). The present amino acid sequence represents a  
 CC clostridial neurotoxin substrate peptide of the invention.

SQ Sequence 116 AA: 100.0%; Score 83; DB 23; Length 116;  
 Query Match Best Local Similarity 100.0%; Pred. No. 8.2e-07;  
 Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 SNKTRIDEANORATKML 17  
 DB 96 SNKTRIDEANORATKML 112  
 RESULT 11  
 AAW30103  
 ID AAW30103 standard; peptide: 206 AA.  
 AC AAW30103;  
 XX  
 XX  
 DT 06-APR-1998 (first entry)  
 DE Synaposomal associated protein.  
 DE Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;  
 KM excitation-secretory uncoupling peptide; catecholamine secretion;  
 KM bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;  
 KM synaposomal associated protein; SNAP-25.  
 KM  
 OS Homo sapiens.  
 XX  
 XX MO9734620-A1.  
 PN 25-SEP-1997.  
 PD  
 XX  
 PF 18-MAR-1997; 97MO-US04393.  
 XX  
 PR 18-MAR-1996; 96US-0013599.  
 XX  
 XX (RECG ) UNIV CALIFORNIA.  
 PA  
 XX  
 PI Montal M;  
 DR  
 XX  
 DR WPI: 1997-479986/44.  
 XX  
 PT Excitation-secretory uncoupling peptide(s) for inhibiting  
 PT neuro-transmitter release - used particularly for treating muscle  
 PT spasticity, and for delivering drugs specifically to neural cells  
 PS  
 XX  
 PS Disclosure: Page 27-28; 61pp: English.  
 CC This sequence represents the human 25 kD synaposomal associated protein  
 CC (SNAP-25), which is an inhibitory agent of the invention. The agents of  
 CC the invention inhibit secretion of neurotransmitter from neuronal cells  
 CC and is an excitation-secretory uncoupling peptide (I) of at least 20  
 CC amino acids (aa) all of which correspond substantially to any one of  
 CC AAW30097-W30102, or more generally any (I) that inhibits 50% of  
 CC catecholamine secretion from bovine chromaffin cells at a concentration  
 CC of 10 microm, especially 0.25 microm, or less. (I) are used, as a  
 CC replacement for Clostridium toxin, to inhibit release of  
 CC neurotransmitters from synaptic vesicles, specifically for reducing  
 CC muscle spasticity. Also (I) may be labeled to allow in vivo imaging of  
 CC intracellular distribution of (I). Compounds for delivering the drug to  
 CC neural cells provide targeted drug delivery, e.g. of substance P to  
 CC brain tumours for induction of apoptosis. Unlike the neurotoxins, (I) are  
 CC not toxic or immunogenic and are more readily available. Their  
 CC therapeutic effect lasts for several days or weeks, so lower doses or  
 CC less frequent treatments are required.  
 CC  
 SQ Sequence 206 AA:  
 Query Match 100.0%; Score 83; DB 18; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 1.6e-06;  
 Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 SNKTRIDEANORATKML 17

DB 187 SNKTRIDEANORATKML 203  
 RESULT 12  
 AAW79198  
 ID AAW79198 standard; Protein: 206 AA.  
 AC AAW79198;  
 XX  
 XX  
 DT 25-NOV-1998 (first entry)  
 DE Mouse SNAP-25 polypeptide.  
 DE Hrs-2 polypeptide; ATP-prefering nucleotidase; SNAP-25; vesicle docking;  
 KM calcium-regulated secretion; secretory vesicle; secretory process; brain;  
 KM neurotransmitter release; presynaptic membrane; CNS disorder; depression;  
 KM Parkinson's disease; endocrine system; hormonal imbalance; cell division;  
 KM thought disorder; schizophrenia; degenerative disorder; anaesthesia; rat;  
 KM immune system; antigen processing; immunomodulator; viral processing;  
 KM central nervous system; vesicular release; affective disorder; human;  
 KM anti-tumour application; membrane trafficking regulation; mouse.  
 OS Mus sp.  
 XX  
 XX MO9838210-A2.  
 PN 03-SEP-1998.  
 PD  
 XX  
 PF 26-FEB-1998; 98MO-US03789.  
 XX  
 PR 26-FEB-1997; 97US-0039159.  
 XX  
 XX (STND ) UNIV LELAND STANFORD JUNIOR.  
 PA  
 XX  
 PI Bean AJ, Scheller RH;  
 DR  
 XX  
 DR WPI: 1998-481140/41.  
 DR N-PSDB; AAV57558.  
 XX  
 PT New isolated Hrs-2 nucleotidase - used in assays to identify  
 PT compounds capable of modulating calcium-regulatory secretion of  
 PT secretory vesicles, such as in neurotransmitter release  
 PS  
 XX  
 PS Claim 16: Pages 42-44; 55pp: English.  
 CC This represents a mouse SNAP-25 polypeptide, a component of the protein  
 CC polypeptides thought to underlie vesicle docking and fusion. The  
 CC invention provides rat and human Hrs-2 polypeptides which are ATP-  
 CC preferring nucleotidase that associate with SNAP-25. For identifying a  
 CC compound capable of modulating calcium-regulated secretion of secretory  
 CC vesicles, a SNAP-25 polypeptide can be contacted with a Hrs-2  
 CC polypeptide. In the presence and absence of a test compound. The effect  
 CC of the test compound on the extent of binding between the SNAP-25 and  
 CC Hrs-2 polypeptides are measured and a compound is identified as effective  
 CC if its measured effect on the extent of binding is above a threshold  
 CC level. The products can be used for identifying drugs capable of  
 CC affecting secretory processes, such as neurotransmitter release at the  
 CC active zones of presynaptic membranes. Such drugs can be used for  
 CC treating disorders or conditions of the central nervous system by  
 CC selectively enhancing or inhibiting vesicular release in specific areas  
 CC of the brain, including affective disorders (e.g. depression), disorders  
 CC of thought (e.g. schizophrenia) and degenerative disorders (Parkinson's  
 CC disease), as well as applications such as anaesthesia. The drugs can  
 CC also be used therapeutically in other systems such as the endocrine  
 CC system for treatment of hormonal imbalances, the immune system for  
 CC intervention in antigen processing, secreted immunomodulators, and viral  
 CC processing, as well as anti-tumour applications, such as regulation of  
 CC membrane trafficking during rapid cell division.  
 CC  
 SQ Sequence 206 AA:  
 Query Match 100.0%; Score 83; DB 19; Length 206;



Best Local Similarity 100.0%; Pred. No. 1.6e-06;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 SNKTRIDEANORATKML 17  
Db 187 SNKTRIDEANORATKML 203

RESULT 13  
AAU00246 ID AAM43426 standard; Protein: 206 AA.  
AAU43426; AAM43426;  
27-APR-1998 (first entry)  
XX Mouse synaptosomal-associated protein-25.  
DE  
XX Binding domain; mouse; syntaxin; synaptosomal-associated protein; CNS;  
KW neurotransmitter; presynaptic membrane; central nervous system; tumour;  
KW neurodegenerative disease; hormonal disorder; immunological disorder.  
XX  
XX Mus sp.  
OS  
XX US5693476-A.  
PN  
XX 02-DEC-1997.  
PD  
XX 24-FEB-1995; 95US-0393985.  
PF  
XX 24-FEB-1995; 95US-0393985.  
PR  
XX (STRD ) UNIV LELAND STANFORD JUNIOR.  
PA  
XX  
PI Scheller RH;  
XX  
XX WPI: 1998-031743/03.  
DR N-PSDB: AAV01554.  
XX  
XX Screening assay for modulators of syntaxin binding - using peptide  
PT for treating binding site of syntaxin, for identifying drugs useful  
PT for treating CNS disorders, neuro-degenerative diseases, etc  
XX  
XX Disclosure; Column 67-72; 57pp; English.  
XX  
XX This amino acid sequence represents the mouse synaptosomal-associated  
CC protein of 25 kd (SNAP-25). The invention relates to a method for  
CC identifying a compound capable of affecting the binding of a  
CC syntaxin-binding protein (SBP), e.g. SNAP-25, alpha-SNAP, n-secl or VAMP,  
CC to syntaxin. The method comprises measuring the effect of the test  
CC compound on the extent of binding between the SBP and the SBP-binding  
CC site on syntaxin. The method can be used for identifying drugs capable  
CC of inhibiting or stimulating neurotransmitter release at the active zones  
CC of presynaptic membranes, which may be useful for treating CNS disorders,  
CC affective or psychotic disorders, neurodegenerative diseases, hormonal or  
CC immunological disorders or tumours.  
XX  
XX Sequence 206 AA;  
SO

Query Match 100.0%; Score 83; DB 19; Length 206;  
Best Local Similarity 100.0%; Pred. No. 1.6e-06;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 SNKTRIDEANORATKML 17  
Db 187 SNKTRIDEANORATKML 203

RESULT 14  
AAU00246 ID AAM00246 standard; Protein: 206 AA.  
AAU00246; AAM00246;

XX 12-SEP-2001 (first entry)  
DT  
XX Synaptosomal-associated protein, SNAP25.  
DE  
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
KW synaptosomal-associated protein; mutagenic; PCR primer; mouse;  
KW N-ethylmaleimide-sensitive fusion protein;  
KW soluble NSF attachment protein receptor.  
XX  
XX Mus sp.  
OS  
XX  
XX  
XX Key Location/Qualifiers  
FH Cleavage-site 180..181  
FT /note= "Peptide bond susceptible to cleavage by  
FT clostridial neurotoxin"  
FT Cleavage-site 197..198  
FT /note= "Peptide bonds susceptible to cleavage by  
FT clostridial neurotoxin"  
FT  
XX WO200118038-A2.  
XX  
XX 15-MAR-2001.  
XX  
XX 18-AUG-2000; 2000WO-GB03196.  
XX  
XX 20-AUG-1999; 99US-0149993.  
XX  
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX  
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
XX  
XX WPI: 2001-226739/23.  
XX  
XX Treating a patient suffering from poisoning or at risk of poisoning by  
PT a clostridial toxin, e.g. botulism, comprises administering a  
PT toxin-resistant or toxin-inhibitory SNARE -  
XX  
XX Disclosure; Fig 8; 131pp; English.  
XX  
XX The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25. The sequence was used to  
CC create SNAP-25 double/single point mutants and C-terminal deletion  
CC mutants used in a new method of treating a patient suffering from  
CC poisoning or at risk of poisoning by a clostridial toxin, comprising  
CC supplying a SNARE (soluble (N-ethylmaleimide-sensitive fusion protein)-  
CC attachment protein receptor) to a cell of the patient, where the SNARE is  
CC resistant to proteolysis by the toxin (toxin-resistant SNARE) and/or is  
CC capable of inhibiting the toxin (toxin-inhibitory SNARE). The protein can  
CC be used in a method of treating a patient in need of inhibition of SNARE-  
CC dependent exocytosis from a cell capable of performing SNARE-dependent  
CC exocytosis, comprises supplying a fragment, variant, fusion or derivative  
CC of a SNARE or an inhibitory SNARE to the cell of the patient. The toxin  
CC resistant or toxin inhibitory SNARE or a recombinant polynucleotide  
CC encoding the SNARE is useful in the manufacture of a medicament for the  
CC treatment of a patient suffering from poisoning or at risk of poisoning  
CC by clostridial toxin, e.g. from botulism or tetanus. The fragment,  
CC variant, fusion or derivative of a SNARE or of an inhibitory SNARE, or a  
CC recombinant polynucleotide encoding either of these SNARE polypeptides  
CC are useful in the manufacture of medicament for the treatment of a  
CC patient in need of inhibition of SNARE-dependent exocytosis from a cell  
CC capable of performing SNARE-dependent exocytosis. The method of treatment  
CC is relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
XX  
XX Sequence 206 AA;  
SO

Query Match 100.0%; Score 83; DB 22; Length 206;  
Best Local Similarity 100.0%; Pred. No. 1.6e-06;  
Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 SNKTRIDEANORATKML 17  
Db 187 SNKTRIDEANORATKML 203

RESULT 15

AAU00252 standard; Protein: 206 AA.

AAU00252:

12-SEP-2001 (first entry)

SNARE homologue, synaptosomal-associated protein, hSNAP25a.

SNAP-25; poisoning: clostridial toxin; SNARE; toxin-resistant SNARE;  
toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
synaptosomal-associated protein; hSNAP25a; human;  
N-ethylmaleimide-sensitive fusion;  
soluble NSF-attachment protein receptor.

Homo sapiens.

WO200118038-A2.

15-MAR-2001.

18-AUG-2000; 2000MO-GB03196.

20-AUG-1999; 99US-0149993.

(IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.

Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;

WPI: 2001-226739/23.

N-PSDB; AAS00369.

Treating a patient suffering from poisoning or at risk of poisoning by  
a clostridial toxin, e.g. botulism, comprises administering a  
toxin-resistant or toxin-inhibitory SNARE -

Disclosure: Fig 8; 131pp; English.

The sequence represents the amino acid sequence of SNARE homologue,  
synaptosomal-associated membrane protein, hSNAP25a, used during analysis  
of SNAP-25. SNAP-25 mutants were used in a new method of treating a  
patient suffering from poisoning or at risk of poisoning by a clostridial  
toxin, comprising supplying a SNARE (soluble (N-ethylmaleimide-sensitive  
fusion protein)-attachment protein receptor) to a cell of the patient,  
where the SNARE is resistant to proteolysis by the toxin (toxin-resistant  
SNARE) and/or is capable of inhibiting the toxin (toxin-inhibitory  
SNARE). The protein can be used in a method of treating a patient in need  
of inhibition of SNARE-dependent exocytosis from a cell capable of  
performing SNARE-dependent exocytosis, comprising supplying a fragment,  
variant, fusion or derivative of a SNARE or an inhibitory SNARE to the  
cell of the patient. The toxin resistant or toxin inhibitory SNARE or a  
recombinant polynucleotide encoding the SNARE is useful in the  
manufacture of a medicament for the treatment of a patient suffering from  
poisoning or at risk of poisoning by clostridial toxin, e.g. from  
botulism or tetanus. The fragment, variant, fusion or derivative of a  
SNARE or of an inhibitory SNARE, or a recombinant polynucleotide encoding  
either of these SNARE polypeptides are useful in the manufacture of  
medicament for the treatment of a patient in need of inhibition of SNARE-  
dependent exocytosis from a cell capable of performing SNARE-dependent  
exocytosis. The method of treatment is relatively fast, thus  
alleviating the symptoms when most severe and taking the patient out of  
critical state.

Sequence 206 AA;

• Query Match 100.0%; Score 83; DB 22; Length 206;  
Best Local Similarity 100.0%; Pred. No. 1.6e-06;

Matches 17; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 SNKTRIDEANORATKML 17  
Db 187 SNKTRIDEANORATKML 203

Search completed: November 19, 2002, 17:39:13  
Job time : 22.3125 secs

	RESULT 1
AA86823	
ID	AA86823 standard; Peptide: 70 AA.
XX	
AC	AA86823;
XX	
D7	15-AUG-1996 (first entry)
XX	
DE	SNAP-25 residues 137-206.
VAMP; vesicle-associated membrane protein; SNAP-25; syntaxin; neurotransmitter; neurotoxin; botulinum; botulism; cleavage; substrate; antibody; detection; assay.	
OS	Synthetic.
PN	W09533850-A1.
PD	14-DEC-1995.
PF	02-JUN-1995; 95WO-GB01279.
PR	03-JUN-1994; 94GB-0011138.
(CAMR-) CAMR CENT APPLIED MICROBIOLOGY & RES. (MICR-) MICROBIOLOGICAL RES AUTHORITY.	
Hallis B, James BAF, Quinn CP, Shone CC; WPI: 1996-040249/04.	
Assay for botulinum or tetanus toxin - by combining test cpd. with substrate which is cleaved by the toxin, and antipod / specific for	

PT the cleaved but not uncleaved substrate  
XX  
PS Example 4: Page 19; 48pp; English.  
XX  
CC The botulinum neurotoxins possess highly specific zinc-endopeptidase  
CC activities within their light sub-units. Depending on the neurotoxin  
CC type these act to cleave small proteins within the nerve cell which are  
CC involved in neurotransmitter release. Antibodies are used in assays  
CC which detect cleaved but not uncleaved substrate. Assays for botulinum  
CC types A and E use the present sequence as a substrate. The sequence is  
CC SNAP-25 protein, residues 137-206.  
XX  
SQ Sequence 70 AA:  
Query Match 100.0%; Score 158; DB 17; Length 70;  
Best Local Similarity 100.0%; Pred. No. 3,1e-16;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 IIGNLRHMLDGMNEIDTQNRQIDRIMEKAD 31  
D6 20 IIGNLRHMLDGMNEIDTQNRQIDRIMEKAD 50  
RESULT 2  
AA015165  
ID AA015165 standard; peptide: 116 AA.  
XX  
AC AA015165;  
XX  
DT 02-SEP-2002 (first entry)  
XX  
DE Clostridial neurotoxin protease substrate peptide 4.  
XX  
KW Clostridial neurotoxin substrate; botulinum neurotoxin substrate; FRET;  
KW fluorescence resonant energy transfer assay; quenched-signal;  
KM clostridial neurotoxin detection; food.  
XX  
OS Unidentified.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 1 /note="S-fluoresceinyl-cysteine"  
FT Cleavage-site 89..90 /note="The peptide is cleaved between these two  
FT residues by type E Clostridium botulinum neurotoxin"  
FT 106..107  
FT /note="The peptide is cleaved between these two  
FT residues by type A Clostridium botulinum neurotoxin"  
XX  
XX WO200225284-A2.  
XX  
XX 28-MAR-2002.  
XX  
XX 25-SEP-2001; 2001WO-US30188.  
XX  
XX 25-SEP-2000; 2000US-235050P.  
XX  
XX (USME-) US MEDICAL RES INST INFECTIOUS DISEASES.  
XX  
XX Schmidt JJ, Stafford RG;  
XX  
XX WPI; 2002-499829/53.  
XX  
XX Substrate useful in e.g. an assay for the protease activity of  
XX clostridial neurotoxin, comprises modified peptide or protein -  
XX  
XX Claim 22; Page 16; 48pp; English.  
XX  
XX The invention comprises clostridial neurotoxin substrate peptides which  
XX can serve as fluorescence resonant energy transfer assay (FRET) or  
XX quenched-signal substrates in assays for the proteolytic activities of  
XX clostridial neurotoxins. The invention further comprises Clostridium  
XX botulinum neurotoxin substrate peptides that can serve as immobilised  
XX clostridial neurotoxins. The invention further comprises Clostridium  
XX botulinum neurotoxin substrate peptides that can serve as immobilised

CC substrates (i.e. bound to a solid phase) in assays for the proteolytic  
CC activities of clostridial neurotoxins. The clostridial (including the  
CC Clostridium botulinum) neurotoxin substrate peptides are useful for  
CC detecting the presence of clostridial neurotoxins in a sample (e.g. food  
CC or an environmental sample). The present amino acid sequence represents a  
CC clostridial neurotoxin substrate peptide of the invention.  
XX  
SQ Sequence 116 AA;  
Query Match 100.0%; Score 158; DB 23; Length 116;  
Best Local Similarity 100.0%; Pred. No. 5,6e-16;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 IIGNLRHMLDGMNEIDTQNRQIDRIMEKAD 31  
D6 65 IIGNLRHMLDGMNEIDTQNRQIDRIMEKAD 95  
RESULT 3  
AA015166  
ID AA015166 standard; peptide: 116 AA.  
XX  
AC AA015166;  
XX  
DT 02-SEP-2002 (first entry)  
XX  
DE Clostridial neurotoxin protease substrate peptide 5.  
XX  
KW Clostridial neurotoxin substrate; botulinum neurotoxin substrate; FRET;  
KW fluorescence resonant energy transfer assay; quenched-signal;  
KM clostridial neurotoxin detection; food.  
XX  
OS Unidentified.  
XX  
FH Key Location/Qualifiers  
FT Modified-site 1 /note="S-fluoresceinyl-cysteine"  
FT Cleavage-site 89..90 /note="The peptide is cleaved between these two  
FT residues by type E Clostridium botulinum neurotoxin"  
XX  
XX WO200225284-A2.  
XX  
XX 28-MAR-2002.  
XX  
XX 25-SEP-2001; 2001WO-US30188.  
XX  
XX 25-SEP-2000; 2000US-235050P.  
XX  
XX (USME-) US MEDICAL RES INST INFECTIOUS DISEASES.  
XX  
XX Schmidt JJ, Stafford RG;  
XX  
XX WPI; 2002-499829/53.  
XX  
XX Substrate useful in e.g. an assay for the protease activity of  
XX clostridial neurotoxin, comprises modified peptide or protein -  
XX  
XX Claim 28; Page 17; 48pp; English.  
XX  
XX The invention comprises clostridial neurotoxin substrate peptides which  
XX can serve as fluorescence resonant energy transfer assay (FRET) or  
XX quenched-signal substrates in assays for the proteolytic activities of  
XX clostridial neurotoxins. The invention further comprises Clostridium  
XX botulinum neurotoxin substrate peptides that can serve as immobilised  
XX substrates (i.e. bound to a solid phase) in assays for the proteolytic  
XX activities of clostridial neurotoxins. The clostridial (including the  
XX Clostridium botulinum) neurotoxin substrate peptides are useful for  
XX detecting the presence of clostridial neurotoxins in a sample (e.g. food  
XX or an environmental sample). The present amino acid sequence represents a  
XX clostridial neurotoxin substrate peptide of the invention.  
XX  
SQ Sequence 116 AA;

```

Query Match          100.0%; Score 158; DB 23; Length 116;
Best Local Similarity 100.0%; Pred. No. 5,6e-16;
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 IIGNLRHMAIDMGNEIDTONROIDRIMEKAD 31
   ||||||||||||||||||||||||||||||||
DB 65 IIGNLRHMAIDMGNEIDTONROIDRIMEKAD 95

RESULT 4
AAU00255
ID AAU00255 standard; Protein; 198 AA.
AC AAU00255;
XX
XX 12-SEP-2001 (first entry)
DE Synaptosomal-associated protein, SNAP25, C-terminal deletion 1-198.
XX
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
XX synaptosomal-associated protein; mouse; mutant; mutein;
XX N-ethylmaleimide-sensitive fusion protein;
XX soluble NSF-attachment protein receptor.
OS Mus sp.
OS Synthetic.
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX
XX 20-AUG-1999; 99US-0149993.
XX
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
XX WPI; 2001-226739/73.
XX
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - ; 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, C-terminal deletion 1-198, used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis
XX from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant
XX polynucleotide encoding either of these SNARE polypeptides are useful in
XX the manufacture of medicament for the treatment of a patient in need of
XX inhibition of SNARE-dependent exocytosis from a cell capable of
XX performing SNARE-dependent exocytosis. The method of treatment is
XX relatively fast, thus alleviating the symptoms when most severe and
XX taking the patient out of critical state.
XX
XX Note: The present sequence is not shown in the specification but is

```

```

CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX
XX Sequence 198 AA:
SQ
XX
XX Query Match          100.0%; Score 158; DB 22; Length 198;
XX Best Local Similarity 100.0%; Pred. No. 1e-15;
XX Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 IIGNLRHMAIDMGNEIDTONROIDRIMEKAD 31
   ||||||||||||||||||||||||||||||||
DB 156 IIGNLRHMAIDMGNEIDTONROIDRIMEKAD 186

RESULT 5
AAU00263
ID AAU00263 standard; Protein; 199 AA.
AC AAU00263;
XX
XX 12-SEP-2001 (first entry)
DE Synaptosomal-associated protein, SNAP25, mutant 1-195(R198T).
XX
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;
XX synaptosomal-associated protein; mouse; mutant; mutein;
XX N-ethylmaleimide-sensitive fusion protein;
XX soluble NSF-attachment protein receptor.
OS Mus sp.
OS Synthetic.
XX
XX Key Location/Qualifiers
XX Misc-difference 198 /note= "Wild-type Arg substituted by Thr"
XX
XX WO200118038-A2.
XX
XX 15-MAR-2001.
XX
XX 18-AUG-2000; 2000WO-GB03196.
XX
XX 20-AUG-1999; 99US-0149993.
XX
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
XX
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
XX WPI; 2001-226739/73.
XX
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1; Page - ; 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant 1-195(R198T), used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis
XX from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant

```

CC polynucleotide encoding either of these SNARE polypeptides are useful in  
CC the manufacture of medicament for the treatment of a patient in need of  
CC inhibition of SNARE-dependent exocytosis from a cell capable of  
CC performing SNARE-dependent exocytosis. The method of treatment is  
CC relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
CC Note: The present sequence is not shown in the specification but is  
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).  
XX  
SQ Sequence 199 AA:  
Query Match 100.0%; Score 158; DB 22; Length 199;  
Best Local Similarity 100.0%; Pred. No. 1e-15;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 IIGNLRHMAIDMGNEIDTQNRQIDRIIMEKAD 31  
DB 156 IIGNLRHMAIDMGNEIDTQNRQIDRIIMEKAD 186  
RESULT 6  
AAU00264 standard; Protein: 200 AA.  
XX  
AC AAU00264;  
XX  
DT 12-SEP-2001 (first entry)  
XX  
DE Synaptosomal-associated protein, SNAP25, mutant 1-200(R198T).  
XX  
KW SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
KW synaptosomal-associated protein; mouse; mutant; muteln;  
KW N-ethylmaleimide-sensitive fusion protein;  
KW soluble NSF-attachment protein receptor.  
XX  
KM  
XX  
OS Mus sp.  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Misc-difference 198 /note= "Wild-type Arg substituted by Thr"  
FT  
XX  
XX WO200118038-A2.  
XX  
XX  
XX PD 15-MAR-2001.  
XX  
XX PF 18-AUG-2000; 2000WO-GB03196.  
XX  
XX PR 20-AUG-1999; 99US-0149993.  
XX  
XX PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX  
XX PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
XX  
XX DR WPI; 2001-226739/23.  
XX  
XX PT Treating a patient suffering from poisoning or at risk of poisoning by  
XX a clostridial toxin, e.g. botulism, comprises administering a  
XX toxin-resistant or toxin-inhibitory SNARE -  
XX  
XX PS Example 1; Page -; 131pp; English.  
XX  
CC The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25, mutant 1-200(R198T), used in a new  
CC method of treating a patient suffering from poisoning or at risk of  
CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
CC from a cell capable of performing SNARE-dependent exocytosis, comprises

CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
CC useful in the manufacture of a medicament for the treatment of a patient  
CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
CC polynucleotide encoding either of these SNARE polypeptides are useful in  
CC the manufacture of medicament for the treatment of a patient in need of  
CC inhibition of SNARE-dependent exocytosis from a cell capable of  
CC performing SNARE-dependent exocytosis. The method of treatment is  
CC relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
CC Note: The present sequence is not shown in the specification but is  
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).  
XX  
SQ Sequence 200 AA:  
Query Match 100.0%; Score 158; DB 22; Length 200;  
Best Local Similarity 100.0%; Pred. No. 1e-15;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 IIGNLRHMAIDMGNEIDTQNRQIDRIIMEKAD 31  
DB 156 IIGNLRHMAIDMGNEIDTQNRQIDRIIMEKAD 186  
RESULT 7  
AAU02637 standard; Protein: 201 AA.  
XX  
AC AAU02637;  
XX  
DT 12-SEP-2001 (first entry)  
XX  
DE Synaptosomal-associated protein, SNAP25, mutant 1-201(R198T).  
XX  
KW SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
KW synaptosomal-associated protein; mouse; mutant; muteln;  
KW N-ethylmaleimide-sensitive fusion protein;  
KW soluble NSF-attachment protein receptor.  
XX  
KM  
XX  
OS Mus sp.  
OS Synthetic.  
XX  
FH Key Location/Qualifiers  
FT Misc-difference 198 /note= "Wild-type Arg substituted by Thr"  
FT  
XX  
XX WO200118038-A2.  
XX  
XX  
XX PD 15-MAR-2001.  
XX  
XX PF 18-AUG-2000; 2000WO-GB03196.  
XX  
XX PR 20-AUG-1999; 99US-0149993.  
XX  
XX PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX  
XX PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
XX  
XX DR WPI; 2001-226739/23.  
XX  
XX PT Treating a patient suffering from poisoning or at risk of poisoning by  
XX a clostridial toxin, e.g. botulism, comprises administering a  
XX toxin-resistant or toxin-inhibitory SNARE -  
XX  
XX PS Example 1; Page -; 131pp; English.  
XX  
CC The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25, mutant 1-201(R198T), used in a new  
CC method of treating a patient suffering from poisoning or at risk of

CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
CC useful in the manufacture of a medicament for the treatment of a patient  
CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
CC polynucleotide encoding either of these SNARE polypeptides are useful in  
CC the manufacture of medicament for the treatment of a patient in need of  
CC inhibition of SNARE-dependent exocytosis. The method of treatment is  
CC performing SNARE-dependent exocytosis. The method of treatment is  
CC relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
CC Note: The present sequence is not shown in the specification but is  
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AA000246).  
XX  
SQ Sequence 201 AA;  
Query Match 100.0%; Score 158; DB 22; Length 201;  
Best Local Similarity 100.0%; Pred. No. 1,1e-15;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
OY 1 IIGNLRHMLDNGNEIDTQNRQIDRIMEKAD 31  
Db 156 IIGNLRHMLDNGNEIDTQNRQIDRIMEKAD 186  
RESULT 8  
AA000265  
ID AA000265 standard; Protein: 202 AA.  
XX  
AC AA000265;  
XX  
DT 12-SEP-2001 (first entry)  
XX  
DE Synaptosomal-associated protein, SNAP25, mutant 1-202(R198T).  
XX  
KM SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
KM synaptosomal-associated protein; mouse; mutant; muteln;  
KM N-ethylmaleimide-sensitive fusion protein;  
KM soluble NSF-attachment protein receptor.  
XX  
OS Mus sp.  
OS Synthetic.  
OS  
FH Key Location/Qualifiers  
FT MISC-difference 198 /note= "Wild-type Arg substituted by Thr"  
XX  
XX W0200118038-A2.  
XX  
XX 15-MAR-2001.  
XX  
XX 18-AUG-2000; 2000WO-GB03196.  
XX  
XX 20-AUG-1999; 99US-0149993.  
XX  
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX  
XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
XX  
XX WPI; 2001-226739/23.  
XX  
XX Treating a patient suffering from poisoning or at risk of poisoning by  
XX a clostridial toxin, e.g. botulism, comprises administering a

PT toxin-resistant or toxin-inhibitory SNARE -  
XX  
XX Example 1; Page - : 131pp; English.  
PS  
XX  
CC The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25, mutant 1-202(R198T), used in a new  
CC method of treating a patient suffering from poisoning or at risk of  
CC poisoning by a clostridial toxin, comprising supplying a SNARE (soluble  
CC (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)  
CC to a cell of the patient, where the SNARE is resistant to proteolysis by  
CC the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the  
CC toxin (toxin-inhibitory SNARE). The protein can be used in a method of  
CC treating a patient in need of inhibition of SNARE-dependent exocytosis  
CC from a cell capable of performing SNARE-dependent exocytosis, comprises  
CC supplying a fragment, variant, fusion or derivative of a SNARE or an  
CC inhibitory SNARE to the cell of the patient. The toxin resistant or toxin  
CC inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is  
CC useful in the manufacture of a medicament for the treatment of a patient  
CC suffering from poisoning or at risk of poisoning by clostridial toxin,  
CC e.g. from botulism or tetanus. The fragment, variant, fusion or  
CC derivative of a SNARE or of an inhibitory SNARE, or a recombinant  
CC polynucleotide encoding either of these SNARE polypeptides are useful in  
CC the manufacture of medicament for the treatment of a patient in need of  
CC inhibition of SNARE-dependent exocytosis. The method of treatment is  
CC performing SNARE-dependent exocytosis. The method of treatment is  
CC relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
CC Note: The present sequence is not shown in the specification but is  
CC derived from the mouse SNAP-25 sequence given in figure 8 (see AA000246).  
XX  
SQ Sequence 202 AA;  
Query Match 100.0%; Score 158; DB 22; Length 202;  
Best Local Similarity 100.0%; Pred. No. 1,1e-15;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 IIGNLRHMLDNGNEIDTQNRQIDRIMEKAD 31  
Db 156 IIGNLRHMLDNGNEIDTQNRQIDRIMEKAD 186  
RESULT 9  
AA002636  
ID AA002636 standard; Protein: 203 AA.  
XX  
AC AA002636;  
XX  
DT 12-SEP-2001 (first entry)  
XX  
DE Synaptosomal-associated protein, SNAP25, mutant 1-203(R198T).  
XX  
KM SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
KM toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
KM synaptosomal-associated protein; mouse; mutant; muteln;  
KM N-ethylmaleimide-sensitive fusion protein;  
KM soluble NSF-attachment protein receptor.  
XX  
OS Mus sp.  
OS Synthetic.  
OS  
FH Key Location/Qualifiers  
FT MISC-difference 198 /note= "Wild-type Arg substituted by Thr"  
XX  
XX W0200118038-A2.  
XX  
XX 15-MAR-2001.  
XX  
XX 18-AUG-2000; 2000WO-GB03196.  
XX  
XX 20-AUG-1999; 99US-0149993.  
XX  
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX  
XX

```

XX Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;
XX
XX WPI: 2001-226739/23.
XX
XX Treating a patient suffering from poisoning or at risk of poisoning by
XX a clostridial toxin, e.g. botulism, comprises administering a
XX toxin-resistant or toxin-inhibitory SNARE -
XX
XX Example 1: Page - : 131pp; English.
XX
XX The sequence represents the amino acid sequence of synaptosomal-
XX associated protein, SNAP25, mutant 1-203(R198T), used in a new
XX method of treating a patient suffering from poisoning or at risk of
XX poisoning by a clostridial toxin, comprising supplying a SNARE (soluble
XX (N-ethylmaleimide-sensitive fusion protein)-attachment protein receptor)
XX to a cell of the patient, where the SNARE is resistant to proteolysis by
XX the toxin (toxin-resistant SNARE) and/or is capable of inhibiting the
XX toxin (toxin-inhibitory SNARE). The protein can be used in a method of
XX treating a patient in need of inhibition of SNARE-dependent exocytosis,
XX comprising from a cell capable of performing SNARE-dependent exocytosis, comprises
XX supplying a fragment, variant, fusion or derivative of a SNARE or an
XX inhibitory SNARE to the cell of the patient. The toxin resistant or toxin
XX inhibitory SNARE or a recombinant polynucleotide encoding the SNARE is
XX useful in the manufacture of a medicament for the treatment of a patient
XX suffering from poisoning or at risk of poisoning by clostridial toxin,
XX e.g. from botulism or tetanus. The fragment, variant, fusion or
XX derivative of a SNARE or of an inhibitory SNARE, or a recombinant
XX polynucleotide encoding either of these SNARE polypeptides are useful in
XX the manufacture of medicament for the treatment of a patient in need of
XX inhibition of SNARE-dependent exocytosis from a cell capable of
XX performing SNARE-dependent exocytosis. The method of treatment is
XX relatively fast, thus alleviating the symptoms when most severe and
XX taking the patient out of critical state.
XX Note: The present sequence is not shown in the specification but is
XX derived from the mouse SNAP-25 sequence given in figure 8 (see AAU00246).
XX
XX
XX Sequence 203 AA.
XX
XX
XX Query Match 100.0%; Score 158; DB 22; Length 203;
XX Best Local Similarity 100.0%; Pred. No. 1,1e-15;
XX Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0
XX
XX 1 IIGNLRHMLDNGNEIDTONRQIDRIEMKAD 31
XX ||||||||||||||||||||||||||||
XX Db 156 IIGNLRHMLDNGNEIDTONRQIDRIEMKAD 186
XX
XX RESULT 10
XX AAW30103
XX ID AAW30103 standard; peptide; 206 AA.
XX
XX AAW30103;
XX
XX 06-APR-1998 (first entry)
XX
XX Synaptosomal associated protein.
XX
XX Neurotransmitter secretion; inhibitor; neuronal cell; synaptic vesicle;
XX excitation-secretory uncoupling peptide; catecholamine secretion;
XX bovine chromaffin cell; Clostridium toxin; muscle spasticity reduction;
XX synaptosomal associated protein; SNAP-25.
XX
XX Homo sapiens.
XX
XX W09734620-A1.
XX
XX 25-SEP-1997.
XX
XX 18-MAR-1997; 97WO-US04393.
XX
XX 18-MAR-1996; 96US-0013559.
XX
XX

```

PA	(RGC ) UNIV CALIFORNIA.
XX	
PI	Montal M;
XX	
DR	WPI: 1997-4/79986/44.
XX	
PT	Excitation-secretory uncoupling peptide(s) for inhibiting
PT	neurotransmitter release - used particularly for treating muscle
PT	spasticity, and for delivering drugs specifically to neural cells
XX	
PS	Disclosure: Page 27-28; 61pp; English.
XX	
CC	This sequence represents the human 25 kD synaptosomal associated protein
CC	(SNAP-25), which is an inhibitory agent of the invention. The agents of
CC	the invention inhibit secretion of neurotransmitter from neuronal cells
CC	and is an excitation-secretory uncoupling peptide (I) of at least 20
CC	amino acids (aa) all of which correspond substantially to any one of
CC	AA330097-330102, or more generally any (I) that inhibits 50% of
CC	catecholamine secretion from bovine chromaffin cells at a concentration
CC	of 10 microm, especially 0.25 microm, or less. (I) are used, as a
CC	replacement for clostridium toxin, to inhibit release of
CC	neurotransmitters from synaptic vesicles, specifically for reducing of
CC	muscle spasticity. Also (I) may be labelled to allow in vivo imaging of
CC	intracellular distribution of (I). Compounds for delivering the drug to
CC	neural cells provide targeted drug delivery, e.g. of substance P to
CC	brain tumours for induction of apoptosis. Unlike the neurotoxins, (I) are
CC	not toxic or immunogenic and are more readily available. Their
CC	therapeutic effect lasts for several days or weeks, so lower doses or
CC	less frequent treatments are required.
XX	
SO	Sequence 206 AA:
Query Match	100.0%; Score 158; DB 18; Length 206;
Best Local Similarity	100.0%; Pred. No. 1,1e-15;
Matches 31; Conservative	0; Mismatches 0; Indels 0; Gaps 0
QY	1 ITGNLRHMLDMGNEIDTNRQIDRIMEXAD 31
Db	156 ITGNLRHMLDMGNEIDTNRQIDRIMEXAD 186
RESULT 11	
AAW79198	
ID	AAW79198 standard; Protein; 206 AA.
XX	
AC	AAW79198;
XX	
DT	25-NOV-1998 (first entry)
XX	
DE	Mouse SNAP-25 polypeptide.
XX	
KW	Hrs-2 polypeptide; ATP-preferring nucleotidase; SNAP-25; vesicle docking;
KW	calcium-regulated secretion; secretory vesicle; secretory process; brain;
KW	neurotransmitter release; presynaptic membrane; CNS disorder; depression;
KW	Parkinson's disease; endocrine system; hormonal imbalance; cell division;
KW	thought disorder; schizophrenia; degenerative disorder; anaesthesia; rat;
KW	immune system; antigen processing; immunomodulator; viral processing;
KW	central nervous system; vesicular release; affective disorder; human;
XX	anti-tumour application; membrane trafficking regulation; mouse.
OS	Mus sp.
XX	
PN	WO9838210-A2.
XX	
PD	03-SEP-1998.
XX	
PF	26-FEB-1998; 98WO-US03789.
XX	
PR	26-FEB-1997; 97US-0039159.
XX	
PA	(STRD ) UNIV LELAND STANFORD JUNIOR.
PI	Bean AJ, Scheller RH;



XX WPI: 1998-481140/41.  
 DR N-PSDB; AAV57558.  
 XX  
 PT New isolated Hrs-2 nucleotidase - used in assays to identify  
 compounds capable of modulating calcium-regulatory secretion of  
 secretory vesicles, such as in neurotransmitter release  
 XX  
 PS Claim 16; Pages 42-44; 55pp; English.  
 XX  
 CC This represents a mouse SNAP-25 polypeptide, a component of the protein  
 CC polypeptides thought to underlie vesicle docking and fusion. The  
 CC invention provides rat and human Hrs-2 polypeptides which are APP-  
 CC preferring nucleotidase that associate with SNAP-25. For identifying a  
 CC compound capable of modulating calcium-regulated secretion of secretory  
 CC vesicles, a SNAP-25 polypeptide can be contacted with a Hrs-2  
 CC polypeptide, in the presence and absence of a test compound. The effect  
 CC of the test compound on the extent of binding between the SNAP-25 and  
 CC Hrs-2 polypeptides are measured and a compound is identified as effective  
 CC if its measured effect on the extent of binding is above a threshold  
 CC level. The products can be used for identifying drugs capable of  
 CC affecting secretory processes, such as neurotransmitter release at the  
 CC active zones of presynaptic membranes. Such drugs can be used for  
 CC treating disorders or conditions of the central nervous system by  
 CC selectively enhancing or inhibiting vesicular release in specific areas  
 CC of the brain, including affective disorders (e.g. depression), disorders  
 CC of thought (e.g. schizophrenia) and degenerative disorders (Parkinson's  
 CC disease), as well as applications such as anaesthesia. The drugs can  
 CC also be used therapeutically in other systems such as the endocrine  
 CC system for treatment of hormonal imbalances, the immune system for  
 CC intervention in antigen processing, secreted immunomodulators, and viral  
 CC processing, as well as anti-tumour applications, such as regulation of  
 CC membrane trafficking during rapid cell division.  
 CC  
 SQ Sequence 206 AA:  
 Query Match 100.0%; Score 158; DB 19; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 1.1e-15;  
 Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 IIGNLRHMLDNGNEIDTQNRQIDRIMERKAD 31  
 DB 156 IIGNLRHMLDNGNEIDTQNRQIDRIMERKAD 186  
 RESULT 12  
 AAW43426  
 ID AAW43426 standard; Protein: 206 AA.  
 AC AAW43426;  
 XX  
 DT 27-APR-1998 (first entry)  
 XX  
 DE Mouse synaptosomal-associated protein-25.  
 XX  
 KW Binding domain; mouse; syntaxin; synaptosomal-associated protein; CNS;  
 KW neurotransmitter; presynaptic membrane; central nervous system; tumour;  
 KW neurodegenerative disease; hormonal disorder; immunological disorder.  
 XX  
 OS Mus sp.  
 PN US5693476-A.  
 PD 02-DEC-1997.  
 PF 24-FEB-1995; 950S-0393985.  
 PR 24-FEB-1995; 950S-0393985.  
 PA (STRD) UNIV LELAND STANFORD JUNIOR.  
 PI Scheller RH;  
 XX

DR WPI: 1998-031743/03.  
 DR N-PSDB; AAV01554.  
 XX  
 PT Screening assay for modulators of syntaxin binding - using peptide  
 PT comprising binding site of syntaxin, for identifying drugs useful  
 PT for treating CNS disorders, neuro-degenerative diseases, etc  
 XX  
 PS Disclosure; Column 67-72; 57pp; English.  
 XX  
 CC This amino acid sequence represents the mouse synaptosomal-associated  
 CC protein of 25 kD (SNAP-25). The invention relates to a method for  
 CC identifying a compound capable of affecting the binding of a  
 CC syntaxin-binding protein (SBP), e.g. SNAP-25, alpha-SNAP, n-secl or VAMP,  
 CC to syntaxin. The method comprises measuring the effect of the test  
 CC compound on the extent of binding between the SBP and the SBP-binding  
 CC site on syntaxin. The method can be used for identifying drugs capable  
 CC of inhibiting or stimulating neurotransmitter release at the active zones  
 CC of presynaptic membranes, which may be useful for treating CNS disorders,  
 CC affective or psychotic disorders, neurodegenerative diseases, hormonal or  
 CC immunological disorders or tumours.  
 CC  
 SQ Sequence 206 AA:  
 Query Match 100.0%; Score 158; DB 19; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 1.1e-15;  
 Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 IIGNLRHMLDNGNEIDTQNRQIDRIMERKAD 31  
 DB 156 IIGNLRHMLDNGNEIDTQNRQIDRIMERKAD 186  
 RESULT 13  
 AAU00246  
 ID AAU00246 standard; Protein: 206 AA.  
 AC AAU00246;  
 XX  
 DT 12-SEP-2001 (first entry)  
 XX  
 DE Synaptosomal-associated protein, SNAP25.  
 XX  
 KW SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
 KW toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
 KW synaptosomal-associated protein; mutagenic; PCR primer; mouse;  
 KW N-ethylmaleimide-sensitive fusion protein;  
 KW soluble NSF-attachment protein receptor.  
 XX  
 OS Mus sp.  
 PN WO200118038-A2.  
 PD 15-MAR-2001.  
 PF 18-AUG-2000; 2000WO-GB03196.  
 PR 20-AUG-1999; 99US-0149993.  
 PA (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
 PI Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
 DR WPI: 2001-226739/23.  
 XX

PT Treating a patient suffering from poisoning or at risk of poisoning by  
PT a clostridial toxin, e.g. botulism, comprises administering a  
PT toxin-resistant or toxin-inhibitory SNARE -  
XX  
PS Disclosure; Fig 8; 131pp; English.  
XX  
CC The sequence represents the amino acid sequence of synaptosomal-  
CC associated protein, SNAP25. The sequence was used to  
CC create SNAP-25 double/single point mutants and C-terminal deletion  
CC mutants used in a new method of treating a patient suffering from  
CC poisoning or at risk of poisoning by a clostridial toxin, comprising  
CC supplying a SNARE (soluble (N-ethylmaleimide-sensitive fusion protein) -  
CC attachment protein receptor) to a cell of the patient, where the SNARE is  
CC resistant to proteolysis by the toxin (toxin-resistant SNARE) and/or is  
CC capable of inhibiting the toxin (toxin-inhibitory SNARE). The protein can  
CC be used in a method of treating a patient in need of inhibition of SNARE-  
CC dependent exocytosis from a cell capable of performing SNARE-dependent  
CC exocytosis, comprises supplying a fragment, variant, fusion or derivative  
CC of a SNARE or an inhibitory SNARE to the cell of the patient. The toxin  
CC resistant or toxin inhibitory SNARE or a recombinant polynucleotide  
CC encoding the SNARE is useful in the manufacture of a medicament for the  
CC treatment of a patient suffering from poisoning or at risk of poisoning  
CC by clostridial toxin, e.g. from botulism or tetanus. The fragment,  
CC variant, fusion or derivative of a SNARE or of an inhibitory SNARE, or a  
CC recombinant polynucleotide encoding either of these SNARE polypeptides  
CC are useful in the manufacture of a medicament for the treatment of a  
CC patient in need of inhibition of SNARE-dependent exocytosis from a cell  
CC capable of performing SNARE-dependent exocytosis. The method of treatment  
CC is relatively fast, thus alleviating the symptoms when most severe and  
CC taking the patient out of critical state.  
XX  
SQ Sequence 206 AA:  
  
Query Match 100.0%; Score 158; DB 22; Length 206;  
Best Local Similarity 100.0%; Pred. No. 1.1e-15;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
OY 1 IIGNLRHMLDNGNEIDTONRQIDRIMEXAD 31  
DB 156 IIGNLRHMLDNGNEIDTONRQIDRIMEXAD 186  
|||||  
RESULT 14  
AAU00252  
ID AAU00252 standard; Protein; 206 AA.  
XX  
AC AAU00252;  
XX  
DT 12-SEP-2001 (first entry)  
XX  
DE SNARE homologue, synaptosomal-associated protein, hSNAP25a.  
XX  
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
XX synaptosomal-associated protein; hSNAP25a; human;  
XX N-ethylmaleimide-sensitive fusion;  
XX soluble NSF-attachment protein receptor.  
XX  
XX Homo sapiens.  
XX  
XX WO200118038-A2.  
XX  
PD 15-MAR-2001.  
XX  
PF 18-AUG-2000; 2000WO-GB03196.  
XX  
PR 20-AUG-1999; 99US-0149993.  
XX  
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX PA Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
XX PI WPI; 2001-226739/23.  
XX  
DR

DR N-PSDB; AAS00369.  
XX  
XX Treating a patient suffering from poisoning or at risk of poisoning by  
PT a clostridial toxin, e.g. botulism, comprises administering a  
PT toxin-resistant or toxin-inhibitory SNARE -  
XX  
PS Disclosure; Fig 8; 131pp; English.  
XX  
XX The sequence represents the amino acid sequence of SNARE homologue,  
CC synaptosomal-associated membrane protein, hSNAP25a, used during analysis  
CC of SNAP-25. SNAP-25 mutants were used in a new method of treating a  
CC patient suffering from poisoning or at risk of poisoning by a clostridial  
CC toxin, comprising supplying a SNARE (soluble (N-ethylmaleimide-sensitive  
CC fusion protein) attachment protein receptor) to a cell of the patient,  
CC where the SNARE is resistant to proteolysis by the toxin (toxin-resistant  
CC SNARE) and/or is capable of inhibiting the toxin (toxin-inhibitory  
CC SNARE). The protein can be used in a method of treating a patient in need  
CC of inhibition of SNARE-dependent exocytosis from a cell capable of  
CC performing SNARE-dependent exocytosis, comprises supplying a fragment,  
CC variant, fusion or derivative of a SNARE or an inhibitory SNARE to the  
CC cell of the patient. The toxin resistant or toxin inhibitory SNARE or a  
CC recombinant polynucleotide encoding the SNARE is useful in the  
CC manufacture of a medicament for the treatment of a patient suffering from  
CC poisoning or at risk of poisoning by clostridial toxin, e.g. from  
CC botulism or tetanus. The fragment, variant, fusion or derivative of a  
CC SNARE or of an inhibitory SNARE, or a recombinant polynucleotide encoding  
CC either of these SNARE polypeptides are useful in the manufacture of  
CC a medicament for the treatment of a patient in need of inhibition of SNARE-  
CC dependent exocytosis from a cell capable of performing SNARE-dependent  
CC exocytosis. The method of treatment is relatively fast, thus  
CC alleviating the symptoms when most severe and taking the patient out of  
CC critical state.  
XX  
SQ Sequence 206 AA:  
  
Query Match 100.0%; Score 158; DB 22; Length 206;  
Best Local Similarity 100.0%; Pred. No. 1.1e-15;  
Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
  
OY 1 IIGNLRHMLDNGNEIDTONRQIDRIMEXAD 31  
DB 156 IIGNLRHMLDNGNEIDTONRQIDRIMEXAD 186  
|||||  
RESULT 15  
AAU00253  
ID AAU00253 standard; Protein; 206 AA.  
XX  
AC AAU00253;  
XX  
DT 12-SEP-2001 (first entry)  
XX  
DE SNARE homologue, synaptosomal-associated protein, hSNAP25b.  
XX  
XX SNAP-25; poisoning; clostridial toxin; SNARE; toxin-resistant SNARE;  
XX toxin-inhibitory SNARE; botulism; tetanus; SNARE-dependent exocytosis;  
XX synaptosomal-associated protein; hSNAP25b; human.  
XX  
XX Homo sapiens.  
XX  
XX WO200118038-A2.  
XX  
PD 15-MAR-2001.  
XX  
PF 18-AUG-2000; 2000WO-GB03196.  
XX  
PR 20-AUG-1999; 99US-0149993.  
XX  
XX (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.  
XX PA Dolly JO, O'Sullivan GA, Mohammed N, Foran PG;  
XX PI WPI; 2001-226739/23.  
XX  
DR

DR N-PSDB; AAS00370.

XX Treating a patient suffering from poisoning or at risk of poisoning by  
 PT a clostridial toxin, e.g. botulism, comprises administering a  
 PT toxin-resistant or toxin-inhibitory SNARE -

XX Disclosure; Fig 8; 130pp; English.

XX The sequence represents the amino acid sequence of SNARE homologue,  
 CC synaptosomal-associated membrane protein, hSNAP25b, used during analysis  
 CC of SNAP-25. SNAP-25 mutants were used in a new method of treating a  
 CC patient suffering from poisoning or at risk of poisoning by a clostridial  
 CC toxin, comprising supplying a SNARE (soluble (N-ethylmaleimide-sensitive  
 CC fusion protein)-attachment protein receptor) to a cell of the patient,  
 CC where the SNARE is resistant to proteolysis by the toxin (toxin-resistant  
 CC SNARE) and/or is capable of inhibiting the toxin (toxin-inhibitory  
 CC SNARE). The protein can be used in a method of treating a patient in need  
 CC of inhibition of SNARE-dependent exocytosis from a cell capable of  
 CC performing SNARE-dependent exocytosis, comprises supplying a fragment,  
 CC variant, fusion or derivative of a SNARE or an inhibitory SNARE to the  
 CC cell of the patient. The toxin resistant or toxin inhibitory SNARE or a  
 CC recombinant polynucleotide encoding the SNARE is useful in the  
 CC manufacture of a medicament for the treatment of a patient suffering from  
 CC poisoning or at risk of poisoning by clostridial toxin, e.g. from  
 CC botulism or tetanus. The fragment, variant, fusion or derivative of a  
 CC SNARE or of an inhibitory SNARE, or a recombinant polynucleotide encoding  
 CC either of these SNARE polypeptides are useful in the manufacture of  
 CC medicament for the treatment of a patient in need of inhibition of SNARE-  
 CC dependent exocytosis from a cell capable of performing SNARE-dependent  
 CC exocytosis. The method of treatment is relatively fast, thus  
 CC alleviating the symptoms when most severe and taking the patient out of  
 CC critical state.

XX Sequence 206 AA:

Query Match 100.0%; Score 158; DB 22; Length 206;  
 Best Local Similarity 100.0%; Pred. No. 1.1e-15;  
 Matches 31; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 ITGNLRHMLDGMGNEDTGNROIDRIMERKAD 31  
 ||||||||||||||||||||||||||||||||  
 DB 156 ITGNLRHMLDGMGNEDTGNROIDRIMERKAD 186

Search completed: November 19, 2002, 17:39:13  
 Job time : 40.6875 secs

**THIS PAGE BLANK (USPTO)**

=> file reg  
FILE 'REGISTRY' ENTERED AT 16:02:48 ON 03 DEC 2002  
USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.  
PLEASE SEE "HELP USAGETERMS" FOR DETAILS.  
COPYRIGHT (C) 2002 American Chemical Society (ACS)

Property values tagged with IC are from the ZIC/VINITI data file  
provided by InfoChem.

STRUCTURE FILE UPDATES: 2 DEC 2002 HIGHEST RN 474876-19-2  
DICTIONARY FILE UPDATES: 2 DEC 2002 HIGHEST RN 474876-19-2

TSCA INFORMATION NOW CURRENT THROUGH MAY 20, 2002

Please note that search-term pricing does apply when  
conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. See HELP  
PROPERTIES for more information. See STNote 27, Searching Properties  
in the CAS Registry File, for complete details:  
<http://www.cas.org/ONLINE/STN/STNOTES/stnotes27.pdf>

=> d que l1  
L1 78 SEA FILE=REGISTRY ABB=ON PLU=ON EANQRA|ANQRAT|NQRATK/SQSP

=> file caplus; d que l7  
FILE 'CAPLUS' ENTERED AT 16:08:29 ON 03 DEC 2002  
USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.  
PLEASE SEE "HELP USAGETERMS" FOR DETAILS.  
COPYRIGHT (C) 2002 AMERICAN CHEMICAL SOCIETY (ACS)

Copyright of the articles to which records in this database refer is  
held by the publishers listed in the PUBLISHER (PB) field (available  
for records published or updated in Chemical Abstracts after December  
26, 1996), unless otherwise indicated in the original publications.  
The CA Lexicon is the copyrighted intellectual property of the  
American Chemical Society and is provided to assist you in searching  
databases on STN. Any dissemination, distribution, copying, or storing  
of this information, without the prior written consent of CAS, is  
strictly prohibited.

FILE COVERS 1907 - 3 Dec 2002 VOL 137 ISS 23  
FILE LAST UPDATED: 2 Dec 2002 (20021202/ED)

This file contains CAS Registry Numbers for easy and accurate  
substance identification.

CAS roles have been modified effective December 16, 2001. Please  
check your SDI profiles to see if they need to be revised. For  
information on CAS roles, enter HELP ROLES at an arrow prompt or use  
the CAS Roles thesaurus (/RL field) in this file.

L1 78 SEA FILE=REGISTRY ABB=ON PLU=ON EANQRA|ANQRAT|NQRATK/SQSP  
L2 44 SEA FILE=CAPLUS ABB=ON PLU=ON L1  
L3 4704 SEA FILE=CAPLUS ABB=ON PLU=ON BOTUL?  
L4 20 SEA FILE=CAPLUS ABB=ON PLU=ON L2 AND L3

THIS PAGE BLANK (USPTO)

L5 1235 SEA FILE=CAPLUS ABB=ON PLU=ON BONT?  
 L6 9 SEA FILE=CAPLUS ABB=ON PLU=ON L2 AND L5  
 L7 20 SEA FILE=CAPLUS ABB=ON PLU=ON L4 OR L6

=> d ibib ab hitrn 17 1-20

L7 ANSWER 1 OF 20 CAPLUS COPYRIGHT 2002 ACS  
 ACCESSION NUMBER: 2002:869489 CAPLUS  
 TITLE: Recombinant light chains of **botulinum**  
 neurotoxins and light chain fusion proteins for use in  
 research and clinical therapy  
 INVENTOR(S): Smith, Leonard; Jensen, Melody  
 PATENT ASSIGNEE(S): USA  
 SOURCE: U.S. Pat. Appl. Publ., 99 pp., Cont.-in-part of U.S.  
 Ser. No. 910,186.  
 CODEN: USXXCO  
 DOCUMENT TYPE: Patent  
 LANGUAGE: English  
 FAMILY ACC. NUM. COUNT: 3  
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 2002168727	A1	20021114	US 2001-11588	20011106
PRIORITY APPLN. INFO.:				
			US 1993-123975	B1 19930921
			US 1999-133865P	P 19990512
			US 1999-133866P	P 19990512
			US 1999-133867P	P 19990512
			US 1999-133868P	P 19990512
			US 1999-133869P	P 19990512
			US 1999-133873P	P 19990512
			US 2000-611419	A1 20000706
			US 2000-246774P	P 20001106
			US 2001-910186	A2 20010720
			US 2001-311966P	P 20010809

AB **Botulinum** neurotoxins, the most potent of all toxins, induce lethal neuromuscular paralysis by inhibiting exocytosis at the neuromuscular junction. The light chains (LC) of these dichain neurotoxins are a new class of zinc-endopeptidases that specifically cleave the synaptosomal proteins, SNAP-25, VAMP, or syntaxin at discrete sites. The present invention relates to the construction, expression, purifn., and use of synthetic or recombinant **botulinum** neurotoxin genes. For example, a synthetic gene for the LC of the **botulinum** neurotoxin serotype A (**BoNT/A**) was constructed and overexpressed in *Escherichia coli*. The gene product was purified from inclusion bodies. The methods of the invention can provide 1.1 g of the LC per L of culture. The LC product was stable in soln. at 4.degree. C. for at least 6 mo. This rBoNT/A LC was proteolytically active, specifically cleaving the Glu-Arg bond in a 17-residue synthetic peptide of SNAP-25, the reported cleavage site of **BoNT/A**. Its calcd. catalytic efficiency  $k_{ub.degree.cat}^{sub.degree.}/K_{ub.degree.m}^{sub.degree.}$  was higher than that reported for the native **BoNT/A** dichain. Treating the rBoNT/A LC with mercuric compds. completely abolished its activity, most probably by modifying the cysteine-164 residue located in the vicinity of the active site. About 70% activity of the LC was restored by adding  $Zn^{up.degree.2+up.degree.}$  to a  $Zn^{up.degree.2+up.degree.}$ -free, apo-LC prepn. The LC was nontoxic to mice and failed to elicit neutralizing epitope(s) when the animals were vaccinated with this protein. In addn., injecting rBoNT/A LC into sea urchin eggs inhibited exocytosis-dependent plasma membrane resealing.

**THIS PAGE BLANK (USPTO)**



IT INDEXING IN PROGRESS

IT 216568-37-5

RL: BSU (Biological study, unclassified); BIOL (Biological study)  
(peptide sybstrate; recombinant light chains of **botulinum**  
neurotoxins and light chain fusion proteins for use in research and  
clin. therapy)

IT 188592-00-9

RL: PRP (Properties)  
(unclaimed sequence; recombinant light chains of **botulinum**  
neurotoxins and light chain fusion proteins for use in research and  
clin. therapy)

L7 ANSWER 2 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2002:595499 CAPLUS

DOCUMENT NUMBER: 137:145554

TITLE: Methods of administering **botulinum** toxin

INVENTOR(S): Walker, Patricia S.

PATENT ASSIGNEE(S): Allergan Sales, Inc., USA

SOURCE: U.S. Pat. Appl. Publ., 33 pp., Cont.-in-part of U. S.  
Ser. No. 730,237.

CODEN: USXXCO

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 2002107199	A1	20020808	US 2002-51952	20020117
US 2002086036	A1	20020704	US 2000-730237	20001205

PRIORITY APPLN. INFO.: US 2000-730237 A2 20001205

AB Methods for treating conditions in an animal or human subject are  
disclosed. The conditions may be pain, skeletal muscle conditions, smooth  
muscle conditions, glandular conditions and cosmetic conditions. The  
methods comprise the step of administering a Clostridium neurotoxin  
component or Clostridium neurotoxin component-encoding DNA to the subject  
using a needleless syringe.

IT 439904-18-4

RL: PRP (Properties)  
(unclaimed sequence; administration of **botulinum** toxin)

L7 ANSWER 3 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2002:505236 CAPLUS

DOCUMENT NUMBER: 137:83622

TITLE: Methods for treating hyperhidrosis

INVENTOR(S): Walker, Patricia S.

PATENT ASSIGNEE(S): Allergan Sales, Inc., USA

SOURCE: U.S. Pat. Appl. Publ., 29 pp.

CODEN: USXXCO

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 2002086036	A1	20020704	US 2000-730237	20001205
US 2002107199	A1	20020808	US 2002-51952	20020117

PRIORITY APPLN. INFO.: US 2000-730237 A2 20001205

AB Methods for treating hyperhidrosis is disclosed herein. In one

**THIS PAGE BLANK (USPTO)**

embodiment, the method includes a step of administering a neurotoxin to a skin area to alleviate excessive sweating. In another embodiment, the method employs a needleless injector to affect the administration of a neurotoxin, for example **botulinum** toxin type A.

IT 439904-18-4

RL: PRP (Properties)

(unclaimed sequence; methods for treating hyperhidrosis)

L7 ANSWER 4 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2002:353597 CAPLUS

DOCUMENT NUMBER: 136:365216

TITLE: Recombinant light chains of **botulinum** neurotoxins and light chain fusion proteins for use in research and clinical therapy

INVENTOR(S): Smith, Leonard A.; Jensen, Melody

PATENT ASSIGNEE(S): United States Army Medical Research and Material Command, USA

SOURCE: PCT Int. Appl., 166 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 3

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2002036758	A2	20020510	WO 2001-US47230	20011106
W:				
AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
RW:				
GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG				
AU 2002028887	A5	20020515	AU 2002-28887	20011106
PRIORITY APPLN. INFO.:			US 2000-246774P	P 20001106
			US 2001-910186	A 20010720
			US 2001-311966P	P 20010809
			WO 2001-US47230	W 20011106

AB **Botulinum** neurotoxins, the most potent of all toxins, induce lethal neuromuscular paralysis by inhibiting exocytosis at the neuromuscular junction. The light chains (LC) of these dichain neurotoxins are a new class of zinc-endopeptidases that specifically cleave the synaptosomal proteins, SNAP-25, VAMP, or syntaxin at discrete sites. The present invention relates to the construction, expression, purifn., and use of synthetic or recombinant **botulinum** neurotoxin genes. For example, a synthetic gene for the LC of the **botulinum** neurotoxin serotype A (**BoNT/A**) was constructed and overexpressed in *Escherichia coli*. The gene product was purified from inclusion bodies. The methods of the invention can provide 1.1 g of the LC per L of culture. The LC product was stable in soln. at 4.degree. for at least 6 mo. This rBoNT/A LC was proteolytically active, specifically cleaving the Glu-Arg bond in a 17-residue synthetic peptide of SNAP-25, the reported cleavage site of **BoNT/A**. Its calcd. catalytic efficiency kcat/Km was higher than that reported for the native **BoNT/A** dichain. Treating the rBoNT/A LC with mercuric compds. completely abolished its activity, most probably by modifying the cysteine-164 residue located in the vicinity of the active site. About 70% activity of the LC was restored by adding Zn2+-free, apo-LC prepn.

**THIS PAGE BLANK (USPTO)**

The LC was nontoxic to mice and failed to elicit neutralizing epitope(s) when the animals were vaccinated with this protein. In addn., injecting rBoNT/A LC into sea urchin eggs inhibited exocytosis-dependent plasma membrane resealing.

IT 216568-37-5

RL: BSU (Biological study, unclassified); BIOL (Biological study) (peptide substrate; recombinant light chains of **botulinum** neurotoxins and light chain fusion proteins for use in research and clin. therapy)

IT 188592-00-9

RL: PRP (Properties) (unclaimed sequence; recombinant light chains of **botulinum** neurotoxins and light chain fusion proteins for use in research and clin. therapy)

L7 ANSWER 5 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2002:286211 CAPLUS

DOCUMENT NUMBER: 136:290338

TITLE: Peptides that mimic the carboxy-terminal domain of SNAP-25 block acetylcholine release at an Aplysia synapse. [Erratum to document cited in CA132:304502]

AUTHOR(S): Aplan, J. P.; Biser, J. A.; Adler, M.; Ferrer-Montiel, A. V.; Montal, M.; Canaves, J. M.; Filbert, M. G.

CORPORATE SOURCE: Neurotoxicology Branch, US Army Medical Research Institute of Chemical Defense, Aberdeen Proving Ground, MD, 21010-5400, USA

SOURCE: Journal of Applied Toxicology (2000), 20(6), 499  
CODEN: JJATDK; ISSN: 0260-437X

PUBLISHER: John Wiley & Sons Ltd.

DOCUMENT TYPE: Journal

LANGUAGE: English

AB The cor. author information is given.

IT 169265-36-5 196928-77-5

RL: ADV (Adverse effect, including toxicity); BIOL (Biological study) (peptides mimicking carboxy-terminal domain of SNAP-25 block acetylcholine release at Aplysia synapse (Erratum))

L7 ANSWER 6 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2002:241093 CAPLUS

DOCUMENT NUMBER: 136:274685

TITLE: Fluorescent substrates and high throughput assays for proteolytic activities of clostridial neurotoxins

INVENTOR(S): Schmidt, James J.; Stafford, Robert G.

PATENT ASSIGNEE(S): U.S. Medical Research Institute of Infectious Diseases, USA

SOURCE: PCT Int. Appl., 48 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2002025284	A2	20020328	WO 2001-US30188	20010925

W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, GM, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NC, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM

**THIS PAGE BLANK (USPTO)**

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,  
DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF,  
BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG

PRIORITY APPLN. INFO.: US 2000-235050P P 20000925

AB In this application is described substrates for high-throughput assays of clostridial neurotoxin proteolytic activities. Two types of substrates are described: (1) modified peptides or proteins that can serve as FRET substrates and (2) modified peptides or proteins that can serve as immobilized substrates. In both types a fluorescent mol. is present in the substrate, eliminating the requirement for the addn. of a fluorogenic reagent. The assays described can be readily adapted for use in automated or robotic systems.

IT 405665-86-3 406458-86-4

RL: ARG (Analytical reagent use); PRP (Properties); ANST (Analytical study); USES (Uses)  
(fluorescent substrates and high throughput assays for proteolytic activities of clostridial neurotoxins)

L7 ANSWER 7 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2001:618890 CAPLUS

DOCUMENT NUMBER: 136:50089

TITLE: High-throughput assays for **botulinum** neurotoxin proteolytic activity: Serotypes A, B, D, and F

AUTHOR(S): Schmidt, James J.; Stafford, Robert G.; Millard, Charles B.

CORPORATE SOURCE: Department of Cell Biology and Biochemistry, Toxicology and Aerobiology Division, U.S. Army Medical Research Institute of Infectious Diseases, Fort Detrick, MD, 21702-5011, USA

SOURCE: Analytical Biochemistry (2001), 296(1), 130-137  
CODEN: ANBCA2; ISSN: 0003-2697

PUBLISHER: Academic Press

DOCUMENT TYPE: Journal

LANGUAGE: English

AB **Botulinum** neurotoxins (**BoNT**) are zinc metalloproteases that cleave and inactivate cellular proteins essential for neurotransmitter release. Because the paralytic effect of **BoNT** is a consequence of its enzymic activity, selective inhibitors may be useful as drugs or as tools for further research. To expedite inhibitor discovery, the authors developed high-throughput, solid-phase protease activity assays for four of the seven **BoNT** serotypes: A, B, D, and F. Each assay consisted of a cleavable oligopeptide, based on the natural substrate sequence, labeled with fluorescein and covalently attached to maleimide-activated multiwell plates. Solns. of holotoxin or nontoxic catalytic domain of **BoNT** were incubated in substrate-coated wells, with or without test compds., followed by transfer and assay of solubilized product in a multiwell fluorometer. Routine toxin concns. ranged from 10 to 100 ng/mL, but concns. as low as 2 ng/mL gave reproducible signals. The fluorescence assays were selective, gave very low background readings, and were stable upon prolonged storage. Using the nontoxic catalytic domain of **BoNT** A, the authors detd. the relative inhibitory potencies of a family of structurally related pseudotripeptide compds. Unlike previous methods, the authors' assays did not employ antibodies or reverse-phase extn. steps, only well-to-well transfers, and were easily adapted to a high-throughput automated environment. (c) 2001 Academic Press.

IT 381670-91-3D, immobilized; fluorescein labeled

RL: ARG (Analytical reagent use); ANST (Analytical study); USES (Uses)  
(substrate; high-throughput solid-phase fluorometric assays for metalloproteinase activities of **botulinum** neurotoxins A, B, D)

**THIS PAGE BLANK (USPTO)**



and F and inhibitory potencies of pseudotripeptides with  
**botulin A)**

REFERENCE COUNT: 36 THERE ARE 36 CITED REFERENCES AVAILABLE FOR THIS  
RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 8 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2001:185784 CAPLUS

DOCUMENT NUMBER: 134:232968

TITLE: Protease-resistant SNARE mutants and the uses thereof  
in rescue of cellular exocytosis for clostridial  
neurotoxin-poisoned patients

INVENTOR(S): Dolly, James Oliver; O'Sullivan, Gregory A.; Mohammed,  
Nadiem; Foran, Patrick G.

PATENT ASSIGNEE(S): Imperial College Innovations Limited, UK

SOURCE: PCT Int. Appl., 131 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2001018038	A2	20010315	WO 2000-GB3196	20000818
WO 2001018038	A3	20011011		

W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN,  
CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR,  
HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,  
LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU,  
SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN,  
YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,  
DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,  
CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG

EP 1210444 A2 20020605 EP 2000-956652 20000818

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,  
IE, SI, LT, LV, FI, RO, MK, CY, AL

PRIORITY APPLN. INFO.: US 1999-149993P P 19990820

WO 2000-GB3196 W 20000818

AB A method of treating a patient suffering from poisoning by clostridial  
toxin wherein a SNARE (sol. (N-ethylmaleimide-sensitive fusion  
protein)-attachment protein receptor) that is resistant to proteolysis by  
the said clostridial toxin (toxin-resistant SNARE) and/or is capable of  
inhibiting the clostridial toxin is supplied to a cell of the patient.  
The SNARE that is resistant to proteolysis may be, synaptosomal-assocd.  
polypeptide of 25 kDa (SNAP-25). The SNAP-25 is preferably resistant to  
proteolysis by **BoNT/A**, **BoNT/E** and **BoNT/C**. A  
method of treating a patient in need of inhibition of SNARE-dependent  
exocytosis from a cell capable of performing SNARE-dependent exocytosis  
wherein a deriv. (inhibitory SNARE) that is capable of inhibiting  
SNARE-dependent exocytosis is supplied to the said cell of the patient.  
The inhibitory SNARE may be a fragment of SNAP-25 that is derivable by  
cleavage of SNAP-25 by **botulinum** toxin A (**BoNT/A**).  
The cell may be, for example, a nerve cell, adreno-chromaffin cell or  
insulin-secreting cell. The SNARE may be supplied to the cell by  
expressing recombinant polynucleotide construct. The SNARE or construct  
may be targeted to a nerve cell, by means of an inactive clostridial  
neurotoxin. The SNARE may be expressed under the target cell-specific  
promoter.

IT 126880-89-5, Protein SNAP 25 (mouse clone p8.52/p8.51  
synaptosome-associated reduced) 154768-88-4 329758-74-9

**THIS PAGE BLANK (USPTO)**

329764-57-0

RL: PRP (Properties)

(unclaimed protein sequence; protease-resistant SNARE mutants and the uses thereof in rescue of cellular exocytosis for clostridial neurotoxin-poisoned patients)

L7 ANSWER 9 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2001:7597 CAPLUS

DOCUMENT NUMBER: 134:91082

TITLE: Peptide inhibitors of neurotransmitter secretion by neuronal cells

INVENTOR(S): Montal, Mauricio; Canaves, Jaume M.; Ferrer-Monteil, Antonio V.

PATENT ASSIGNEE(S): The Regents of the University of California, USA

SOURCE: U.S., 23 pp.

CODEN: USXXAM

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
-----	---	-----	-----	-----
US 6169074	B1	20010102	US 1997-819286	19970318
PRIORITY APPLN. INFO.:			US 1996-13599P	P 19960318

AB The invention consists of peptides which inhibit the secretion of neurotransmitters from synaptic vesicles. The peptides of the invention are believed to mimic the activity of neurotoxins produced by *Clostridium botulinum* and tetani (including *botulinum* serotypes A, B, C, D, E, F and G). Structurally, the peptides are comprised of amino acid fragments from the substrate binding domains selected from three proteins which bind to form a receptor for docking of synaptic vesicles to the plasma membranes of neuronal cells; i.e., SNAP-25, VAMP-2 and syntaxin. Certain of the inventive peptides exhibit strong inhibitory activity; e.g., 50% or greater decline in neurotransmitter release is obtained at even nanomolar concns. The peptides are suited for use as substitutes for *Clostridium* neurotoxins in clin. applications and in compds. for targeted delivery of drugs into neural cells.

IT 169265-36-5 196928-77-5 197099-52-8

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

(*Clostridium* neurotoxin-mimicking peptide inhibitors of neurotransmitter secretion by neuronal cells)

IT 126880-89-5, Protein SNAP 25 (mouse clone p8.52/p8.51 synaptosome-associated reduced)

RL: PRP (Properties)

(unclaimed protein sequence; peptide inhibitors of neurotransmitter secretion by neuronal cells)

REFERENCE COUNT: 7 THERE ARE 7 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 10 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 2000:132673 CAPLUS

DOCUMENT NUMBER: 132:304502

TITLE: Peptides that mimic the carboxy-terminal domain of SNAP-25 block acetylcholine release at an Aplysia synapse

AUTHOR(S): Apland, J. P.; Biser, J. A.; Adler, M.; Ferrer-Montiel, A. V.; Montal, M.; Filbert, M. G.

CORPORATE SOURCE: Neurotoxicology Branch, US Army Medical Research

**THIS PAGE BLANK (USPTO)**

SOURCE: Institute of Chemical Defense, Aberdeen Proving Ground, MD, 21010-5400, USA  
Journal of Applied Toxicology (1999), 19(Suppl. 1), S23-S26  
CODEN: JJATDK; ISSN: 0260-437X

PUBLISHER: John Wiley & Sons Ltd.

DOCUMENT TYPE: Journal

LANGUAGE: English

AB **Botulinum** neurotoxin serotypes A and E (**BoNT/A** and **BoNT/E**) block neurotransmitter release, presumably by cleaving SNAP-25, a protein involved in docking of synaptic vesicles with the presynaptic plasma membrane. Three excitation-secretion uncoupling peptides (ESUPs), which mimic the carboxy-terminal domain of SNAP-25 and span or adjoin the cleavage sites for **BoNT/A** and **BoNT/E**, also inhibit transmitter release from permeabilized bovine chromaffin cells. In this study, these peptides were tested for effects on acetylcholine (ACh) release at an identified cholinergic synapse in isolated buccal ganglia of *Aplysia californica*. The presynaptic neuron was stimulated elec. to elicit action potentials. The postsynaptic neuron was voltage-clamped, and evoked inhibitory postsynaptic currents (IPSCs) were recorded. The ESUPs were pressure-injected into the presynaptic neuron, and their effects on the amplitude of the IPSCs were studied. Acetylcholine release from presynaptic cells, as measured by IPSC amplitudes, was gradually inhibited by the ESUPs. All three peptides caused .apprx.40% redn. in IPSC amplitude in 2 h. Random-sequence peptides of the same amino acid compn. had no effect. Injection of **BoNT/E**, in contrast, caused .apprx.50% redn. in IPSC amplitude in 30 min and almost complete inhibition in 2 h. These results are the first demonstration that ESUPs block neuronal cholinergic synaptic transmission. They are consistent with the concept that ESUPs compete with the intact SNAP-25 for binding with other fusion proteins, thus inhibiting stimulus-evoked exocytosis of neurotransmitter.

IT 169265-36-5 196928-77-5  
RL: ADV (Adverse effect, including toxicity); BIOL (Biological study) (peptides that mimic carboxy-terminal domain of SNAP-25 block acetylcholine release at *Aplysia* synapse)

REFERENCE COUNT: 16 THERE ARE 16 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 11 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1999:655992 CAPLUS

DOCUMENT NUMBER: 131:268976

TITLE: Assay for the proteolytic activity of **botulin** neurotoxin type A from *Clostridium botulinum*, substrate requirements and activation by serum albumin

INVENTOR(S): Schmidt, James J.; Bostian, Karen A.

PATENT ASSIGNEE(S): United States Dept. of the Army, USA

SOURCE: U.S., 28 pp.  
CODEN: USXXAM

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

PATENT INFORMATION:

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
	US 5965699	A	19991012	US 1996-743894	19961106
AB	A label-based assay is described, through modifications of peptide substrate structure and derivatization of serum albumin, which can be used to det. proteolytic activity of <b>botulin</b> neurotoxin A (botox A)				

THIS PAGE BLANK (USPTO)

without sepn. of products. The present invention provides a method for screening compds. for botox A inhibitory or stimulatory activity. Substrate requirements for botox A were also studied.

IT 245360-92-3 245360-93-4 245360-95-6  
245360-96-7 245360-97-8 245360-98-9  
245360-99-0 245361-00-6 245361-01-7  
245361-02-8 245361-03-9 245361-04-0  
245361-05-1 245361-09-5 245361-11-9  
245361-13-1 245361-16-4 245361-18-6

RL: ARG (Analytical reagent use); ANST (Analytical study); USES (Uses)  
(assay for proteolytic activity of **botulin** neurotoxin A from *Clostridium botulinum*, substrate requirements and activation by serum albumin)

IT 188591-98-2 188591-99-3 188592-01-0  
188592-02-1 188592-03-2 188592-04-3  
188592-05-4 188592-16-7 188592-17-8  
188592-18-9 245361-21-1

RL: BPR (Biological process); BSU (Biological study, unclassified); BIOL (Biological study); PROC (Process)  
(assay for proteolytic activity of **botulin** neurotoxin A from *Clostridium botulinum*, substrate requirements and activation by serum albumin)

REFERENCE COUNT: 4 THERE ARE 4 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 12 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1999:636059 CAPLUS

DOCUMENT NUMBER: 131:268231

TITLE: Antibody-based assay for **botulinum** and tetanus neurotoxins

INVENTOR(S): Shone, Clifford Charles; Hallis, Bassam; James, Benjamin Arthur Frederick; Quinn, Conrad Pdraig

PATENT ASSIGNEE(S): Microbiological Research Authority, UK

SOURCE: U.S., 21 pp., Cont.-in-part of Appl. No. PCT/GB95/01279.

CODEN: USXXAM

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 5962637	A	19991005	US 1996-760001	19961203
WO 9533850	A1	19951214	WO 1995-GB1279	19950602
W: AU, CA, JP, US				
RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
US 6043042	A	20000328	US 1998-15960	19980130
US 6337386	B1	20020108	US 2000-534572	20000327
PRIORITY APPLN. INFO.:			GB 1994-11138	A 19940603
			WO 1995-GB1279	A2 19950602
			US 1996-760001	A3 19961203
			US 1998-15960	A1 19980130

AB The invention provides an antibody-based assay for toxins having peptidase activity, and in particular, this invention relates to assays for **botulinum** and tetanus neurotoxins. The invention comprises the steps of: (a) combining a test compd. with a substrate and with antibody, wherein the substrate has a cleavage site for the toxin and when cleaved by toxin forms a product, and wherein the antibody binds to the product but not to the substrate; and (b) testing for the presence of antibody bound to the product, which product is attached to a solid phase assay

**THIS PAGE BLANK (USPTO)**



component. Preferably, the substrate is a peptide or a protein which is cleaved by the toxin to generate new peptides have N- and C-terminal ends. In addn., the target peptide is preferably selected from the group VAMP, SNAP-25, and syntaxin, and it may also be from analogs, isoforms, and/or fragments thereof. Furthermore, the assay is capable of distinguishing between active and inactive toxin present within the sample, since inactive toxin will have reduced or no activity.

IT 173080-83-6

RL: PRP (Properties)

(unclaimed protein sequence; antibody-based assay for **botulinum** and tetanus neurotoxins)

REFERENCE COUNT: 8 THERE ARE 8 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 13 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1998:630583 CAPLUS

DOCUMENT NUMBER: 130:21584

TITLE: The 26-mer peptide released from SNAP-25 cleavage by **botulinum** neurotoxin E inhibits vesicle docking

AUTHOR(S): Ferrer-Montiel, Antonio V.; Gutierrez, Luis M.; Apland, James P.; Canaves, Jaume M.; Gil, Anabel; Viniegra, Salvador; Biser, Jennifer A.; Adler, Michael; Montal, Mauricio

CORPORATE SOURCE: Department of Biology, University of California San Diego, La Jolla, CA, 92093-0366, USA

SOURCE: FEBS Letters (1998), 435(1), 84-88

CODEN: FEBLAL; ISSN: 0014-5793

PUBLISHER: Elsevier Science B.V.

DOCUMENT TYPE: Journal

LANGUAGE: English

AB **Botulinum** neurotoxin E (**BoNT** E) cleaves SNAP-25 at the C-terminal domain releasing a 26-mer peptide. This peptide product may act as an excitation-secretion uncoupling peptide (ESUP) to inhibit vesicle fusion and thus contribute to the efficacy of **BoNT** E in disabling neurosecretion. We have addressed this question using a synthetic 26-mer peptide which mimics the amino acid sequence of the naturally released peptide, and is hereafter denoted as ESUP E. This synthetic peptide is a potent inhibitor of Ca<sup>2+</sup>-evoked exocytosis in permeabilized chromaffin cells and reduces neurotransmitter release from identified cholinergic synapses in in vitro buccal ganglia of *Aplysia californica*. In chromaffin cells, both ESUP E and **BoNT** E abrogate the slow component of secretion without affecting the fast, Ca<sup>2+</sup>-mediated fusion event. Anal. of immunoppts. of the synaptic ternary complex involving SNAP-25, VAMP and syntaxin demonstrates that ESUP E interferes with the assembly of the docking complex. Thus, the efficacy of **BoNTs** as inhibitors of neurosecretion may arise from the synergistic action of cleaving the substrate and releasing peptide products that disable the fusion process by blocking specific steps of the exocytotic cascade.

IT 196928-77-5

RL: BAC (Biological activity or effector, except adverse); BPR (Biological process); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study); PROC (Process)

(SNAP-25 peptide ESUP E; 26-mer peptide released from SNAP-25 cleavage by **botulinum** neurotoxin E which inhibits synaptic vesicle docking)

REFERENCE COUNT: 30 THERE ARE 30 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 14 OF 20 CAPLUS COPYRIGHT 2002 ACS

**THIS PAGE BLANK USPTO**

ACCESSION NUMBER: 1998:630507 CAPLUS  
DOCUMENT NUMBER: 130:34875  
TITLE: Type A **botulinum** neurotoxin proteolytic activity: development of competitive inhibitors and implications for substrate specificity at the S1' binding-subsite  
AUTHOR(S): Schmidt, James J.; Stafford, Robert G.; Bostian, Karen A.  
CORPORATE SOURCE: Toxinology Division, United States Army Medical Research Institute of Infectious Diseases, Fort Detrick, MD, 21702-5011, USA  
SOURCE: FEBS Letters (1998), 435(1), 61-64  
CODEN: FEBLAL; ISSN: 0014-5793  
PUBLISHER: Elsevier Science B.V.  
DOCUMENT TYPE: Journal  
LANGUAGE: English

AB Type A **botulinum** neurotoxin (botox A) is a zinc metalloprotease that cleaves only one peptide bond in the synaptosomal protein, SNAP-25. Single-residue changes in a 17-residue substrate peptide were used to develop the first specific, competitive inhibitors of its proteolytic activity. Substrate analog peptides with P4, P3, P2' or P3' cysteine were readily hydrolyzed by the toxin, but those with P1 or P2 cysteine were not cleaved and were inhibitors. Peptides with either D- or L-cysteine as the N-terminus, followed by the last six residues of the substrate, were the most effective inhibitors, each with a Ki value of 2 .mu.M. Elimination of the cysteine sulfhydryl group yielded much less effective inhibitors, suggesting that inhibition was primarily due to binding of the active-site zinc by the sulfhydryl group. Botox A displayed an unusual requirement for arginine as the P1' inhibitor residue, demonstrating that the S1' binding subsite of botox A is dissimilar to those of most other zinc metalloproteases. This characteristic is an important element in shaping the substrate specificity of botox A.

IT 216568-37-5 216568-38-6 216568-39-7  
216568-46-6

RL: BPR (Biological process); BSU (Biological study, unclassified); BIOL (Biological study); PROC (Process)  
(inhibitors and substrates of botox A, a type A **botulinum** neurotoxin with proteolytic activity)

REFERENCE COUNT: 20 THERE ARE 20 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 15 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1997:640559 CAPLUS  
DOCUMENT NUMBER: 127:298730  
TITLE: Peptide neurotoxin analog inhibitors of neurotransmitter secretion by neuronal cells for neural targeting of drugs  
INVENTOR(S): Montal, Mauricio  
PATENT ASSIGNEE(S): Regents of the University of California, USA  
SOURCE: PCT Int. Appl., 61 pp.  
CODEN: PIXXD2  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
FAMILY ACC. NUM. COUNT: 2  
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9734620	A1	19970925	WO 1997 US4392	19970319
W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, GH, HU, IL, IS, JP, KE, KG, KP, KR, KZ,				

**THIS PAGE BLANK (USPTO)**

W 19970318

**THIS PAGE BLANK (USPTO)**

(IPSCs) were recorded. ESUP was pressure-injected into the presynaptic neuron, and its effect on the amplitude of the IPSCs was studied. ACh release from presynaptic cells, as measured by the amplitudes of IPSCs, was consistently inhibited. The inhibition was gradual, requiring 1-3 h to effect a 50-60% redn. of IPSC amplitude. A random-sequence peptide of the same amino acid compn. had no effect. Apparently, ESUP competes with the intact SNAP-25 for binding with other fusion proteins, thus inhibiting exocytosis of neurotransmitter. This effect may account, in part, for **botulinum** toxin-induced inhibition of transmitter release.

IT 169265-36-5, Excitation-secretion uncoupling peptide  
RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); BIOL (Biological study)  
(blockade of acetylcholine release at synapse in Aplysia by peptide that mimics C-terminal domain of SNAP-25)

L7 ANSWER 17 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1997:152896 CAPLUS  
DOCUMENT NUMBER: 126:235112  
TITLE: Endoproteinase activity of type A **botulinum** neurotoxin: substrate requirements and activation by serum albumin  
AUTHOR(S): Schmidt, James J.; Bostian, Karen A.  
CORPORATE SOURCE: Toxinology Division, U.S. Army Medical Res. Institute Infectious Diseases, Frederick, MD, 21702-5011, USA  
SOURCE: Journal of Protein Chemistry (1997), 16(1), 19-26  
CODEN: JPCHD2; ISSN: 0277-8033  
PUBLISHER: Plenum  
DOCUMENT TYPE: Journal  
LANGUAGE: English

AB Type A **botulinum** neurotoxin, a zinc-dependent endoproteinase that selectively cleaves the neuronal protein SNAP-25, can also cleave relatively short peptides. We found that bovine and other serum albumins stimulated the type A-catalyzed hydrolysis of synthetic peptide substrates, through a direct effect on the kinetic consts. of the reaction. Furthermore, with bovine serum albumin in the assays, the optimum substrate size was 16 residues (11 on the amino-terminal side of the cleavage site and 5 on the carboxy-terminal side). To further investigate the catalytic requirements of the neurotoxin, peptides were synthesized with various amino acid substitutions at the P5 through P5' substrate sites. Changes at all of these locations affected values for both kcat and Km. Substitutions at the P2, P1', and P2' sites had more pronounced effects on hydrolysis rates than did substitutions at the P1 site. Enzyme-substrate interactions at the P3' threonine probably involved the side-chain Me group rather than the hydroxyl group. Replacing the P2' alanine with leucine eliminated detectable hydrolysis, but not binding, since this peptide was an inhibitor. A neg. charged residue was preferred at P5, but not at P4. The data indicate that type A **botulinum** neurotoxin has an extended substrate recognition region and a requirement for arginine as the P1' residue.

IT 188591-98-2 188591-99-3 188592-00-9  
188592-01-0 188592-02-1 188592-03-2  
188592-04-3 188592-05-4 188592-16-7  
188592-17-8 188592-18-9  
RL: BPR (Biological process); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study); PROC (Process)  
(endoproteinase activity of type A **botulinum** neurotoxin, substrate requirements and activation by serum albumin)

L7 ANSWER 18 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1997:97334 CAPLUS  
DOCUMENT NUMBER: 126:197791

**THIS PAGE BLANK (USPTO)**



TITLE: A peptide that mimics the C-terminal sequence of SNAP-25 inhibits secretory vesicle docking in chromaffin cells

AUTHOR(S): Gutierrez, Luis M.; Viniegra, Salvador; Rueda, Joaquin; Ferrer-Montiel, Antonio V.; Canaves, Jaume M.; Montal, Mauricio

CORPORATE SOURCE: Instituto de Neurociencias and Facultad de Medicina, Universidad de Alicante, Alicante, 03080, Spain

SOURCE: Journal of Biological Chemistry (1997), 272(5), 2634-2639

CODEN: JBCHA3; ISSN: 0021-9258

PUBLISHER: American Society for Biochemistry and Molecular Biology

DOCUMENT TYPE: Journal

LANGUAGE: English

AB Excitation-secretion uncoupling peptides (ESUPs) are inhibitors of Ca<sup>2+</sup>-dependent exocytosis in neural and endocrine cells. Their mechanism of action, however, remains elusive. We report that ESUP-A, a 20-mer peptide patterned after the C terminus of SNAP-25 (synaptosomal associated protein of 25 kDa) and containing the cleavage sequence for **botulinum** neurotoxin A (**BoNT A**), abrogates the slow, ATP-dependent component of the exocytotic pathway, without affecting the fast, ATP-independent, Ca<sup>2+</sup>-mediated fusion event. Ultrastructural analysis indicates that ESUP-A induces a drastic accumulation of dense-core vesicles near the plasma membrane, mimicking the effect of **BoNT A**. Together, these findings argue in favor of the notion that ESUP-A inhibits ATP-primed exocytosis by blocking vesicle docking. Identification of blocking peptides which mimic sequences that bind to complementary partner domains on interacting proteins of the exocytotic machinery provides new pharmacological tools to dissect the molecular and mechanistic details of neurosecretion. Our findings may assist in developing ESUPs as substitute drugs to **BoNTs** for the treatment of spasmodic disorders.

IT 169265-36-5

RL: BAC (Biological activity or effector, except adverse); BSU (Biological study, unclassified); PRP (Properties); BIOL (Biological study) (amino acid sequence of; SNAP-25 C-terminal sequence-like peptide ESUP-A inhibits ATP-dependent secretory vesicle docking in bovine adrenal chromaffin cells)

L7 ANSWER 19 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1996:87093 CAPLUS

DOCUMENT NUMBER: 124:109558

TITLE: Toxin assay

INVENTOR(S): Shone, Clifford Charles; Hallis, Bassam; James, Benjamin Arthur Frederick; Quinn, Conrad Padraig

PATENT ASSIGNEE(S): Microbiological Research Authority, UK

SOURCE: PCT Int. Appl., 47 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 2

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9533850	A1	19951214	WO 1995-GB1279	19950602
W: AU, CA, JP, US				
RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
AU 9526240	A1	19960104	AU 1995-26240	19950602
AU 687564	B2	19980226		

**THIS PAGE BLANK (0870)**

EP 763131 A1 19970319 EP 1995-921033 19950602  
EP 763131 B1 19990825  
R: AT, BE, CH, DE, DK, FR, GB, IT, LI, NL, SE  
JP 10504801 T2 19980512 JP 1995-500544 19950602  
AT 183779 E 19990915 AT 1995-921033 19950602  
US 5962637 A 19991005 US 1996-760001 19961203  
US 6043042 A 20000328 US 1998-15960 19980130  
US 6337386 B1 20020108 US 2000-534572 20000327  
PRIORITY APPLN. INFO.: GB 1994-11138 A 19940603  
WO 1995-GB1279 W 19950602  
US 1996-760001 A3 19961203  
US 1998-15960 A1 19980130

AB A toxin assay that uses a substrate for cleavage by the toxin and antibodies that do not recognize the substrate but recognize and bind to the product of cleavage of the substrate by the toxin. The substrate can be a nerve cell peptide when the assay is for **botulinum** toxin or tetanus toxin.

IT **173080-83-6**  
RL: ARG (Analytical reagent use); ANST (Analytical study); USES (Uses) (substrate; tetanus and **botulinum** toxin assay using peptide substrates and antibodies)

L7 ANSWER 20 OF 20 CAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1996:19380 CAPLUS

DOCUMENT NUMBER: 124:167929

TITLE: Proteolysis of synthetic peptides by type A **botulinum** neurotoxin

AUTHOR(S): Schmidt, James J.; Bostian, Karen A.

CORPORATE SOURCE: U.S. Army Medical Res. Inst. of Infectious Diseases, Fort Detrick, Frederick, MD, 21702-5011, USA

SOURCE: Journal of Protein Chemistry (1995), 14(8), 703-8  
CODEN: JPCHD2; ISSN: 0277-8033

PUBLISHER: Plenum

DOCUMENT TYPE: Journal

LANGUAGE: English

AB Type A **botulinum** neurotoxin catalyzed the hydrolysis of synthetic peptides based on the sequence of the 25 kDa synaptosomal protein SNAP-25. In each peptide, the toxin cleaved at a single glutaminyl-arginine bond corresponding to residues 197 and 198 of SNAP-25, confirming earlier reports on the enzymic specificity of the toxin in synaptosomal preps. Metal chelators inhibited catalysis, consistent with a metalloprotease activity. In contrast to tetanus toxin and other **botulinum** toxin serotypes, type A toxin hydrolyzed relatively short, 17-to 20-residue peptides. In the substrates, SNAP-25 residue 202 and one or more of residues 197-191 were required for efficient hydrolysis, but residues 167-186 and 203-206 were not. The highest rates of hydrolysis were found when the C-terminal residues of the peptides were amidated.

IT **169265-36-5 172486-01-0 172486-02-1**

**172486-03-2 172486-04-3 172486-05-4**

**172486-06-5 173762-25-9**

RL: BPR (Biological process); BSU (Biological study, unclassified); BIOL (Biological study); PROC (Process)

(proteolysis of synthetic peptides by type A **botulinum** neurotoxin)

=> file home

FILE 'HOME' ENTERED AT 16:11:07 ON 03 DEC 2002

**THIS PAGE BLANK (8870)**